GenCore version 5.1.3
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OM protein - nucleic search, using frame_plus_p2n model

Run on: January 12, 2003, 10:11:31 ; Search time 52 Seconds
                (without alignments)
                1710.313 Million cell updates/sec

Title: US-09-649-108-1
Perfect score: 1511
Sequence: 1 MRIFAVFIFMTYWHLLNAFT..........KCGIQDTNSKKQSDTHLEET 290

Scoring table: BLOSUM62
                Xgapop 10.0 , Xgapext 0.5
                Ygapop 10.0 , Ygapext 0.5
                Fgapop   6.0 , Fgapext   7.0
                Delop   6.0 , Delext   7.0

/ched: 441362 seqs, 153338381 residues
                Maximum Match 100%
                listing first 45 summaries

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%

Command line parameters:
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-Q=/cgn2_1/USPTO_spool/US09649108/runat_12012003_101121_13898/app.query.fasta_1.455
-DB=Issued_Patents_NA -QFMT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cd1
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
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-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database :   Issued_Patents_NA:*
                1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
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                3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
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                5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	311	20.6	1602	4	US-09-651-200-11	Sequence 11, Appl
2	311	20.6	2229	4	US-09-651-200-5	Sequence 5, Appli
3	307	20.3	1020	4	US-09-651-200-7	Sequence 7, Appli
4	307	20.3	1323	4	US-09-651-200-9	Sequence 9, Appli
5	307	20.3	2691	4	US-09-651-200-1	Sequence 1, Appli
6	307	20.3	2885	4	US-09-651-200-3	Sequence 3, Appli
7	191	12.6	2627	4	US-09-404-879A-391	Sequence 391, App
8	176.5	11.7	1491	2	US-08-147-772-1	Sequence 1, Appli
9	176.5	11.7	1491	2	US-08-456-104-5	Sequence 5, Appli
10	176.5	11.7	1491	2	US-08-101-624-22	Sequence 22, Appli
11	176.5	11.7	1491	2	US-08-751-767A-5	Sequence 5, Appli
12	176.5	11.7	1491	3	US-08-153-262-1	Sequence 1, Appli

RESULT 1
US-09-651-200-11

Sequence 11, Application US/09651200
Patent No. 6429303

GENERAL INFORMATION:
APPLICANT: Green et al
TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 15966-562 (CURA-62)
CURRENT APPLICATION NUMBER: US/09/651,200
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/152283
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/172909
PRIOR FILING DATE: 1999-12-21
PRIOR APPLICATION NUMBER: 60/183578
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 11
LENGTH: 1602
TYPE: DNA
ORGANISM: Homo sapiens
US-09-651-200-11

Alignment Scores:
Pred. No.:   4.76e-30
Score: 311.00
Percent Similarity: 48.03%
Best Local Similarity: 30.11%
Query Match: 20.58%
DB: 4

US-09-649-108-1 (1-290) x US-09-651-200-11 (1-1602)

ALIGNMENTS

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Conservative:	50					
Mismatches:	131					
Indels:	14					
Gaps:	5					

13	176.5	11.7	1491	3	US-08-479-744A-28	Sequence 28, Appl
14	176.5	11.7	1491	3	US-08-280-757B-28	Sequence 28, Appli
15	176.5	11.7	1491	3	US-09-159-135-1	Sequence 1, Appli
16	176.5	11.7	1491	4	US-08-205-697A-18	Sequence 18, Appli
17	176.5	11.7	1491	4	US-08-702-525-18	Sequence 18, Appl
18	176.5	11.7	1491	4	US-09-450-798-1	Sequence 1, Appli
19	176.5	11.7	1491	4	US-09-326-1B6B-225	Sequence 225, App
20	176.5	11.7	1491	4	US-08-403-253A-1	Sequence 1, Appli
21	176.5	11.7	1491	2	PCT-US95-02576-18	Sequence 18, Appl
22	176.5	11.7	867	2	US-08-184-009-207	Sequence 207, App
23	172	11.4	867	2	US-08-458-356-207	Sequence 207, App
24	172	11.4	867	2	US-08-812-946A-2	Sequence 2, Appli
25	172	11.4	867	4	US-08-460-736-207	Sequence 207, App
26	172	11.4	879	4	US-09-039-982A-31	Sequence 31, Appli
27	172	11.4	879	4	US-09-039-641-31	Sequence 31, Appli
28	172	11.4	879	4	US-09-039-762A-31	Sequence 31, Appli
29	172	11.4	879	4	US-09-042-492D-31	Sequence 31, Appli
30	172	11.4	879	4	US-08-913-612A-31	Sequence 31, Appli
31	169.5	11.2	921	4	US-08-009-202	Sequence 202, App
32	169.5	11.2	921	4	US-08-356-202	Sequence 202, App
33	169.5	11.2	921	4	US-08-460-736-202	Sequence 202, Appl
34	169.5	11.2	1716	2	US-08-147-772-3	Sequence 3, Appli
35	169.5	11.2	1716	2	US-08-456-104-7	Sequence 7, Appli
36	169.5	11.2	1716	2	US-08-101-624-24	Sequence 24, Appli
37	169.5	11.2	1716	2	US-08-155-262-3	Sequence 3, Appli
38	169.5	11.2	1716	3	US-08-479-744A-30	Sequence 30, Appli
39	169.5	11.2	1716	3	US-08-280-757B-30	Sequence 30, Appli
40	169.5	11.2	1716	3	US-09-159-135-3	Sequence 3, Appli
41	169.5	11.2	1716	4	US-08-205-697A-16	Sequence 16, Appli
42	169.5	11.2	1716	4	US-08-702-525-16	Sequence 16, Appli
43	169.5	11.2	1716	4	US-09-450-798-3	Sequence 3, Appli
44	169.5	11.2	1716	5	PCT-US95-02576-16	Sequence 16, Appli
45	169.5	11.2	1888	4	US-08-205-697A-1	Sequence 1, Appli

; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 5
; LENGTH: 2229
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (60)..(1661)
DB: US-09-651-200-5

US-09-649-108-1 (1-290) x US-09-651-200-5 (1-2229)

Alignment Scores:
Pred. No.:
Score: 311.00   Length: 2229
Percent Similarity: 48.03%   Matches: 84
Best Local Similarity: 30.11%   Conservative: 50
Query Match: 20.58%   Mismatches: 131
                                        Indels: 14
                                        Gaps: 5

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Db   795  GCCGTGGAGGTCCAGGTCCTGAGGACCCGGTGGGCCCCTAGTGGGCACCGGCAATGCCACC  795

Qy    38  IleGluCysLysPheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleValTyrTrp  57
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Db   795  GCCGTGGAGGTCCAGGTCCTGAGGACCCGGTGGGCCCCTAGTGGGCACCGGCAATGCCACC  854

Qy    58  GluMetGluAspProLysAsnIleIleGlnPheValHisGlyGluGluAspLeuLysValGln  77
          |||   || :|||  :||  |||| |||||||||| |||
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Qy    78  HisSerSerTyrArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAla  97
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Qy    98  AlaLeuGlnIleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSer  117
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RESULT 2
US-09-651-200-5
Sequence 5, Application US/09651200
Patent No. 6429303

OM protein - protein search, using sw model

GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.

Run on: January 12, 2003, 10:10:41

Title: US-09-649-108-1_COPY_30_290
Perfect score: 1356
Sequence: 1 VEYGSNMTIECKFPVEKQLD..........KCGIQDTNSKKQSDTHLEET 261

Scoring table: BLOSUM62
                  Gapop 10.0 , Gapext 0.5
Searched: 262574 seqs, 29422922 residues

Post-processing: Minimum Match 0%
                  Maximum Match 100%
                  Listing first 45 summaries

Database :   Issued_Patents_AA:*
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                  4: /cgn2_6/ptodata/2/1aa/6B_COMB.pep:*
                  5: /cgn2_6/ptodata/2/1aa/PCTUS_COMB.pep:*
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000

number of hits satisfying chosen parameters: 262574

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

Search time 14.4076 Seconds
(without alignments)
533.008 Million cell updates/sec

SUMMARIES

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2	292	21.5	534	4	US-08-479-744A-29	Sequence 29, Appli
3	288	21.2	534	4	US-08-280-757B-29	Sequence 29, Appli
4	288	21.2	340	4	US-09-651-200-2	Sequence 2, Appli
5	187.5	13.8	441	4	US-09-651-200-2	Sequence 2, Appli
6	187.5	13.8	282	4	US-09-404-879A-393	Sequence 393, Appli
7	181	13.3	309	4	US-09-404-879A-392	Sequence 392, App
8	177.5	13.1	329	4	US-09-651-200-19	Sequence 19, Appl
9	177.5	13.1	306	2	US-08-147-772-4	Sequence 4, Appli
10	177.5	13.1	306	2	US-08-456-104-8	Sequence 8, Appli
11	177.5	13.1	306	3	US-09-651-624-25	Sequence 25, Appli
12	177.5	13.1	306	4	US-08-479-744A-31	Sequence 31, Appl
13	177.5	13.1	306	4	US-08-280-757B-31	Sequence 31, Appl
14	177.5	13.1	306	3	US-09-159-135-4	Sequence 4, Appli
15	170	12.5	288	4	US-09-450-798-4	Sequence 4, Appli
16	169.5	12.5	306	4	US-09-651-200-14	Sequence 14, Appl
17	169.5	12.5	306	4	US-08-205-697A-17	Sequence 17, Appl
18	169.5	12.5	306	4	US-08-702-525-17	Sequence 17, Appl
19	169.5	12.5	306	4	US-09-651-525-2	Sequence 2, Appl
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23	169.5	12.5	320	5	PCT-US95-02576-2	Sequence 2, Appli
24	167	12.3	288	2	US-08-147-772-2	Sequence 2, Appli
25	167	12.3	288	2	US-08-456-104-2	Sequence 2, Appli
26	167	12.3	288	2	US-08-101-624-23	Sequence 23, Appli
27	167	12.3	288	2	US-08-751-767A-6	Sequence 6, Appli
28	167	12.3	288	3	US-08-153-262-2	Sequence 2, Appli
29	167	12.3	288	4	US-08-479-744A-29	Sequence 29, Appli
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31	167	12.3	288	4	US-09-651-200-2	Sequence 2, Appli
32	167	12.3	288	4	US-08-205-697A-19	Sequence 19, Appli
33	167	12.3	288	4	US-08-702-525-19	Sequence 19, Appl
34	167	12.3	288	4	US-09-450-798-2	Sequence 2, Appli
35	167	12.3	288	4	US-08-403-253A-2	Sequence 2, Appli
36	167	12.3	288	4	US-09-651-200-13	Sequence 13, Appl
37	167	12.3	288	5	PCT-US95-02576-19	Sequence 19, Appli
38	167	12.3	323	5	PCT-US94-09642-2	Sequence 2, Appli
39	161	11.9	323	5	US-09-651-200-21	Sequence 21, Appli
40	161	11.9	329	2	US-08-456-104-2	Sequence 2, Appli
41	161	11.9	329	2	US-08-101-624-2	Sequence 2, Appli
42	161	11.9	329	3	US-08-479-744A-2	Sequence 2, Appli
43	161	11.9	329	4	US-08-280-757B-2	Sequence 2, Appli
44	161	11.9	329	4	US-08-205-697A-23	Sequence 23, Appli
45	161	11.9	329	4	US-08-702-525-23	Sequence 23, Appli

ALIGNMENTS

RESULT 1
US-09-651-200-6
Sequence 6, Application US/09651200
Patent No. 6429303
GENERAL INFORMATION:
APPLICANT: Green et al
TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 15966-562 (CURA-62)
CURRENT APPLICATION NUMBER: US/09/651,200
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/152383
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/172909
PRIOR FILING DATE: 1999-12-21
PRIOR APPLICATION NUMBER: 60/183578
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 6
LENGTH: 534
TYPE: PRT
ORGANISM: Homo sapiens

Query Match   21.5%; Score 292; DB 4; Length 534;
Best Local Similarity   29.5%; Pred. No. 1.7e-22;
Matches   78; Conservative 49; Mismatches 123; Indels 14; Gaps 5;

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Db    261 GTDADLRCSFSPEPGSFSLAQLNLIWQLTDTK--QLVHSFTEGRDQGSSAYANRTALFPDLL 318

Qy     64 SLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNK------INQRILVVDP 117
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Db    319 AQGNASLRLQRVRVADEGSFTCFVSIRDFGSAAVSLQVAAPYSKPSMTLEPNKDLRPGDT 378

Qy    118 VTSEHELTCQA-EGYPKAEVIWTSSDHQVLSGKTTTNSKREEKLFNVTSTLRINTTTNE 176
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Db    379 VT----ITCSSYRGYPEAEVFWQDGGGVPLIGNVTTSQMANEQGLFDVHSVLRVLGANG 434

Qy    177 IFYCTFRRLDPEENHTAELVIPELPLAHPPNERTHVILGAILLCLGVALTFIFRLRKGR 236
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RESULT 2
US-09-651-200-24
Sequence 24, Application US/09651200
Patent No. 6429303
GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: Patentin Ver. 2.0
;
; SEQ ID NO 24
; LENGTH: 534
; TYPE: PRT
; ORGANISM: Unknown
; FEATURE:
; OTHER INFORMATION: Description of Unknown Organism: Sequence
; OTHER INFORMATION: mz5020.protein from Figure 4.
US-09-651-200-24

Query Match   78%; Score 292; DB 4; Length 534;
Best Local Similarity   21.5%; Pred. No. 1.7e-22;
Matches   78; Conservative   49; Mismatches 123; Indels 14; Gaps   5;

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Db  261 GTDATLRCFSFSPEPGFSLAQLNLIWQLTDTK--QLVHSFTEGRDQGSAYANRTALFPDLL 318

Qy   64 SLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNK-----INQRILVVDP 117
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Db  319 AQGNASLRLQRVRVADEGSFTCFVSIRDFGSAAVSLQVAAPYSKPSMTLEPNKDLRPGDT 378

Qy  118 VTSEHELTCQA-EGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNE 176
        || ||| |  |  ||| ||| | |     |  |    | :| ||||:|| ||| :
Db  379 VT----ITCSSYRGYPEAEVFWQDGQGVPLTGNVTTSQMANEQGLFDVHSVLRVVLGANG 434

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Db  495 SCEEENAGAEDQDGEGEGSKTALQ 518
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RESULT 3
US-09-651-200-2
Sequence 2, Application US/09651200
Patent No. 6429303
GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: Patentin Ver. 2.0
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; SEQ ID NO 2
; LENGTH: 340
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-200-2

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Best Local Similarity   21.2%; Pred. No. 2.3e-22;
Matches   77; Conservative   49; Mismatches 124; Indels 14; Gaps   5;

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Qy   64 SLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNK-----INQRILVVDP 117
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Qy  118 VTSEHELTCQA-EGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNE 176
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Db  301 SCEEENAGAEDQDGEGEGSKTALQ 324
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RESULT 4
US-09-651-200-4
Sequence 4, Application US/09651200
Patent No. 6429303
GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: Patentin Ver. 2.0
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; SEQ ID NO 4
; LENGTH: 441
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-200-4

Query Match   77%; Score 288; DB 4; Length 441;
Best Local Similarity   21.2%; Pred. No. 3.3e-22;
Matches   77; Conservative   49; Mismatches 124; Indels 14; Gaps   5;

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Qy   64 SLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNK-----INQRILVVDP 117
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Qy  118 VTSEHELTCQA-EGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNE 176
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Db
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OM protein - protein search, using sw model

Copyright (c) 1993 - 2003 Compugen Ltd.   GenCore version 5.1.3

Run on: January 12, 2003, 10:10:41 ; Search time 11.5924 Seconds
    533.008 Million cell updates/sec

Title: US-09-649-108-10

Perfect score: 1096

Sequence: 1 VEYGSNMTIECKFPVEKQLD..........HTAELVIPELPLAHPNPERT 210
     (without alignments)

Scoring table: BLOSUM62

    Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

l number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
    Maximum Match 100%
    Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	Match DB	ID	Description
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2	265.5	24.2	534	4	US-09-651-200-24	Sequence 24, Appl
3	261.5	23.9	340	4	US-09-651-200-2	Sequence 2, Appl
4	261.5	23.9	441	4	US-09-651-200-2	Sequence 2, Appl
5	179	16.3	282	4	US-09-404-879A-393	Sequence 393, Appl
6	179	16.3	309	4	US-09-404-879A-392	Sequence 392, App
7	164	15.0	288	4	US-09-651-200-14	Sequence 14, Appl
8	159	14.5	306	2	US-08-147-772-4	Sequence 4, Appl
9	159	14.5	306	2	US-08-456-104-8	Sequence 8, Appl
10	159	14.5	306	3	US-08-101-624-25	Sequence 25, Appl
11	159	14.5	306	3	US-08-153-262-4	Sequence 4, Appl
12	159	14.5	306	4	US-08-479-744A-31	Sequence 31, Appl
13	159	14.5	306	4	US-08-280-757B-31	Sequence 31, Appl
14	159	14.5	306	4	US-09-450-798-4	Sequence 4, Appl
15	159	14.5	306	4	US-09-159-135-4	Sequence 4, Appl
16	155	14.1	288	2	US-08-147-772-2	Sequence 2, Appl
17	155	14.1	288	2	US-08-456-104-6	Sequence 6, Appl
18	155	14.1	288	2	US-08-751-767-6	Sequence 6, Appl
19	155	14.1	288	3	US-08-101-624-23	Sequence 23, Appl
20	155	14.1	288	3	US-08-153-262-2	Sequence 2, Appl
21	155	14.1	288	3	US-08-479-744A-29	Sequence 29, Appl
22	155	14.1	288	4	US-08-280-757B-29	Sequence 29, Appl
23	155	14.1	288	4	US-09-159-135-2	Sequence 2, Appl
24	155	14.1	288	4	US-08-205-697A-19	Sequence 19, Appl
25	155	14.1	288	4	US-09-702-525-19	Sequence 19, Appl
26	155	14.1	288	4	US-09-450-798-2	Sequence 2, Appl
27	155	14.1	288	4	US-09-403-253A-2	Sequence 2, Appl
28	155	14.1	288	4	US-09-651-200-13	Sequence 13, Appl
29	155	14.1	288	5	PCT-US95-02576-19	Sequence 19, Appl
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31	153.5	14.0	208	3	US-08-630-172-15	Sequence 15, Appl
32	153.5	14.0	208	4	US-09-375-419-15	Sequence 15, Appl
33	151.5	13.8	208	4	US-09-460-384-36	Sequence 36, Appl
34	151.5	13.8	473	4	US-09-171-945-131	Sequence 131, App
35	151.5	13.8	306	4	US-09-651-200-15	Sequence 15, Appl
36	151	13.8	306	4	US-08-205-697A-17	Sequence 17, Appl
37	151	13.8	306	5	PCT-US95-02576-17	Sequence 17, Appl
38	151	13.8	306	4	US-08-205-525-17	Sequence 17, Appl
39	151	13.8	320	5	PCT-US95-02576-17	Sequence 17, Appl
40	151	13.8	320	5	US-08-702-525-2	Sequence 2, Appl
41	151	13.8	320	4	PCT-US95-02576-2	Sequence 2, Appl
42	151	13.8	329	4	US-09-651-200-18	Sequence 18, Appl
43	149	13.6	299	4	US-09-651-200-15	Sequence 15, Appl
44	144.5	13.2	589	2	US-08-724-39A-1	Sequence 1, Appl
45	140.5	12.8	292	4	US-09-651-200-16	Sequence 16, Appl

ALIGNMENTS

RESULT 1
US-09-651-200-6
Sequence 6, Application US/09651200
; Patent NO. 6429303
; GENERAL INFORMATION:
; APPLICANT: Green et al
; TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
; TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
; TITLE OF INVENTION: Polypeptides Encoded Thereby
; FILE REFERENCE: 15966-562 (CURA-62)
; CURRENT APPLICATION NUMBER: US/09/651,200
; CURRENT FILING DATE: 2000-08-30
; PRIOR APPLICATION NUMBER: 60/152383
; PRIOR FILING DATE: 1999-09-03
; PRIOR APPLICATION NUMBER: 60/172909
; PRIOR FILING DATE: 1999-12-21
; PRIOR APPLICATION NUMBER: 60/183578
; PRIOR FILING DATE: 2000-02-18
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 6
; LENGTH: 534
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-200-6

Query Match   24.2%;   Score 265.5;   DB 4;   Length 534;
Best Local Similarity   31.0%;   Pred. No. 4.5e-20;
Matches   65;   Conservative   38;   Mismatches   94;   Indels   13;   Gaps   4;

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Db   261   GTDATLRCSFSPEPGFSLAQLNLIWQLTDTK---QLVHSFTEGRDQGSSAYANRTALFPDLL   318

Qy   64   SLGNAALQITDVKLQQDAGVYRCMISYGGADYKRITVKVNAPYNK------INQRILVVDP   117
      : |||:||: ||:| || ::|||| |||:|| |:|
Db   319   AQGNASLRLQRVRVADEGSFTCFVSIRDFGSAAVSLQVAAPYSKPSMTLEPNKDLRPGDT   378

Qy   118   VTSEHELTCQA-EGYPKAEVIWTSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNE   176
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Db   379   VT---ITCSSYRGYPEAEVFWQDGQGVPLTGNVTTSQMANEQGLFDVHSVLRVVLGANG   434

Qy   177   IFYCTFRRLDPEEENHTAELVIPELPLAHPP   206
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RESULT 2
US-09-651-200-24
; Sequence 24, Application US/09651200

Patent No. 6429303
GENERAL INFORMATION:
APPLICANT: Green et al
TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 15966-562 (CURA-62)
CURRENT APPLICATION NUMBER: US/09/651,200
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/152383
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/172909
PRIOR FILING DATE: 1999-12-21
PRIOR APPLICATION NUMBER: 60/183578
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 24
LENGTH: 534
TYPE: PRT
ORGANISM: Unknown
FEATURE:
OTHER INFORMATION: Description of Unknown Organism: Sequence
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US-09-651-200-24

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Query Match          24.2%;   Score 265.5;   DB 4;   Length 534;
Best Local Similarity  31.0%;   Pred. No. 4.5e-20;
Matches  65;   Conservative  38;   Mismatches  94;   Indels  13;   Gaps  4;

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Qy  64 SLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNK------INQRILVVDP 117
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RESULT 3
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Patent No. 6429303
GENERAL INFORMATION:
APPLICANT: Green et al
TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 15966-562 (CURA-62)
CURRENT APPLICATION NUMBER: US/09/651,200
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/152383
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/172909
PRIOR FILING DATE: 1999-12-21
PRIOR APPLICATION NUMBER: 60/183578
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 340
TYPE: PRT
ORGANISM: Homo sapiens
US-09-651-200-2

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Best Local Similarity  30.5%;   Pred. No. 6.4e-20;
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RESULT 4
US-09-651-200-4
Sequence 4, Application US/09651200
Patent No. 6429303
GENERAL INFORMATION:
APPLICANT: Green et al
TITLE OF INVENTION: Polynucleotides Encoding Members of the Human B
TITLE OF INVENTION: Lymphocyte Activation Antigen B-7 Family and
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 15966-562 (CURA-62)
CURRENT APPLICATION NUMBER: US/09/651,200
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/152383
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/172909
PRIOR FILING DATE: 1999-12-21
PRIOR APPLICATION NUMBER: 60/183578
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
LENGTH: 441
TYPE: PRT
ORGANISM: Homo sapiens
US-09-651-200-4

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Best Local Similarity  30.5%;   Pred. No. 9.2e-20;
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RESULT 5
US-09-404-879A-393
Sequence 393, Application US/0404879A
Patent No. 6448546
GENERAL INFORMATION:

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.

Run on: January 12, 2003, 11:11:11 ; Search time 62 seconds
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Title: US-09-649-108-1
Perfect score: 1511
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result NO.	Score	Query Match Length	% DB	DB ID	Description
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2	1511	100.0	1553	12	US-10-002-775-3	Sequence 3, Appli
3	1511	100.0	1604	10	US-09-875-338-1	Sequence 1, Appli
4	1511	100.0	3575	10	US-09-796-858-41	Sequence 41, Appli

RESULT 1
US-09-910-174A-22

Alignment Scores:
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Score: 1511.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%

Length: 873
Matches: 290
Conservative: 0
Mismatches: 0

ALIGNMENTS

	Score		DB	ID	
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7	1134	75.0	10	US-09-875-338-4	Sequence 4, Appli
8	1050	69.5	12	US-10-002-775-10	Sequence 10, Appli
9	925.5	61.3	9	US-09-796-858-43	Sequence 43, Appli
10	726	48.0	10	US-09-867-701-10728	Sequence 10728, A
11	713	47.2	10	US-09-867-701-2957	Sequence 2957, Ap
12	565	37.4	10	US-09-867-701-3638	Sequence 18, Appl
13	527	34.9	10	US-09-875-338-18	Sequence 18, Appl
14	442	29.2	10	US-09-895-866-1	Sequence 3, Appli
15	497	32.9	9	US-09-955-866-1	Sequence 3, Appli
16	464	30.7	10	US-09-794-210-1	Sequence 1, Appli
17	891	10	US-09-896-913A-3		Sequence 20, Appli
18	842	10	US-09-875-338-20		Sequence 20, Appl
19	819	10	US-09-895-837-3		Sequence 1, Appli
20	819	10	US-09-895-866-1		Sequence 1, Appli
21	819	10	US-09-896-913A-1		Sequence 1, Appli
22	819	10	US-09-895-837-1		Sequence 1, Appli
23	819	10	US-09-910-174A-1		Sequence 14, Appl
24	842	10	US-09-875-338-14		Sequence 14, Appl
25	1655	9	US-09-796-858-16		Sequence 16, Appl
26	1655	10	US-09-794-210-3		Sequence 4, Appli
27	1655	10	US-09-895-837-4		Sequence 3, Appli
28	741	9	US-09-895-913A-6		Sequence 4, Appli
29	741	10	US-09-895-837-6		Sequence 6, Appli
30	744	10	US-09-794-210-5		Sequence 5, Appli
31	744	10	US-09-910-174A-30		Sequence 30, Appli
32	744	10	US-09-875-338-10		Sequence 10, Appli
33	951	10	US-09-875-338-12		Sequence 12, Appl
34	951	10	US-09-910-174A-23		Sequence 23, Appl
35	1517	9	US-09-790-622-2		Sequence 2, Appli
36	1517	10	US-09-789-561-63		Sequence 63, Appl
37	1998	9	US-09-978-295A-136		Sequence 136, App
38	1998	9	US-09-978-697-136		Sequence 136, App
39	1998	9	US-09-978-192A-136		Sequence 136, App
40	1998	9	US-09-999-832A-136		Sequence 136, App
41	1998	9	US-09-978-189-136		Sequence 136, App
42	1998	12	US-10-052-586-53		Sequence 53, Appl
43	3197	10	US-09-875-338-19		Sequence 19, Appl
44	3197	10	US-09-875-338-19		Sequence 19, Appl
45	951	9	US-09-915-789A-2		Sequence 2, Appli

Query Match: Indels: 0
DB: 100.00%   Gaps: 0
                 10

US-09-649-108-1 (1-290) x US-09-910-174A-22 (1-873)

Qy   1 MetArgIlePheAlaValPheIlePheMetThrTyrThrPheHisLeuLeuAsnAlaPheThr 20
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 ATGAGGATATTTGCTGTGTTTTATATTCATGACCTACTGGCATTTACT 60

Qy   21 ValThrValProLysAspLeuTyrValValGlyTyrGlySerAsnMetThrIleGluCys 40
         ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 GTCACGGTTCCCAAGGACCTATATGTGGTAGGATATGGCAGCAATATGAAATGC 120

Qy   41 LysPheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleValIleTyrArgTrpGluMetGlu 60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 121 AAATTCCCAGTGGGAAAAACAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAG 180

Qy   61 AspLysAsnIleIleGlnPheValHisGlyGluGluAspLeuLysValGlnHisSerSer 80
         ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 181 GATAAGAACATTATTCAATTTGTGCATGGAGAGGAAGACCTGAAGGTTCAGCATAGTAGC 240

Qy   81 TyrArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGln 100
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 241 TACAGACAGAGGGCCCGGCTGTTGAAGGACCAGCTCTCCCTGGGAAATGCTGCACTTCAG 300

Qy 101 IleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerTyrGlyGly 120
         |||||||||||||||||||||||||||||||||||||||||||||||||||
Db 301 ATCACAGATGTGAAATTGCAGGATGCAGGGGTGTACCGCTGCATGATCAGCTATGGTGGT 360

Qy 121 AlaAspTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg 140
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 361 GCCGACTACAAGCGAATTACTGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGA 420

Qy 141 IleLeuLeuValAlaAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr 160
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 421 ATTTTGCTTGTGGTTGGGGATCCAGTCACCTCTGAACATGAACTGACCTGCCAGGCTGAGGGCTAC 480

Qy 161 ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThrThr 180
         |||||||||||||||||||||||||||||||||||||||||||||||||||
Db 481 CCCAAGGCCGAAGTCATCTGGACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCAC 540

Qy 181 ThrThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIleAsn 200
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 541 ACCACCAATTCCAAGAGAAGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATCAAC 600

Qy 201 ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsnHis 220
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 601 ACAACAACTAATGAGATTTTCTACTGCACTTTAGGGAGATTAGATCCTGAGGAAAACCAT 660

Qy 221 ThrAlaGluLeuValIleIleProGluLeuProLeuAlaHisProProAsnGluArgThrHis 240
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 661 ACAGCTGAATTGGTCATCCCAGAACTACCTCTGGCACATCCTCCAAATGAAAGGACTCAC 720

Qy 241 LeuValIleIleLeuGlyAlaIleLeuLeuCysLeuGlyValAlaLeuThrPheIlePheArg 260
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 721 TTGGTAATTCTGGGAGCCATCTTATTATGCCTTGGTGTAGCCCTGACACTTCACCTTCCGT 780

Qy 261 LeuArgLysGlyArgMetMetAspValLysLysCysGlyIleGlnAspThrAsnSerLys 280
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 781 TTAAGAAAAGGGAGAATGATGGATGTGAAAAAATGTGGCATCCAAGATACAAACTCAAAG 840

Qy 281 LysGlnSerAspThrHisLeuGluGluThr 290
         ||||||||||||||||||||||||
Db 841 AAGCAAAGTGATCACACATTTGGAGGAGACG 870

RESULT 2
US-10-002-775-3

; US-10-002-775-3
; Sequence 3, Application US/10002775
; Patent No. US20020102651A1
; GENERAL INFORMATION:
;   APPLICANT:
;   APPLICANT: Gordon Freeman
;   APPLICANT: Vassiliki Boussiotis
;   APPLICANT: Tatyana Chernova

Alignment Scores:
Pred. No.:
Score: 1.27e-174   Length: 1553
Percent Similarity: 1511.00   Matches: 290
Best Local Similarity: 100.00%   Conservative: 0
Query Match: 100.00%   Mismatches: 0
DB: Indels: 0
                                       Gaps: 0

US-09-649-108-1 (1-290) x US-10-002-775-3 (1-1553)

Qy   1 MetArgIlePheAlaValPheIlePheMetThrTyrThrPheHisLeuLeuAsnAlaPheThr 20
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   53 ATGAGGATATTTGCTGTGTTTTATATTCATGACCTACTGGCATTTACT 112

Qy   21 ValThrValProLysAspLeuTyrValValGlyTyrGlySerAsnMetThrIleGluCys 40
         |||||||||||||||||||||||||||||||||||||||||||||||||
Db 113 GTCACGGTTCCCAAGGACCTATATGTGGTAGGATATGGCAGCAATATGAAATGC 172

Qy   41 LysPheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleValIleTyrArgTrpGluMetGlu 60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 173 AAATTCCCAGTGGGAAAAACAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAG 232

Qy   61 AspLysAsnIleIleGlnPheValHisGlyGluGluAspLeuLysValGlnHisSerSer 80
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 233 GATAAGAACATTATTCAATTTGTGCATGGAGAGGAAGACCTGAAGGTTCAGCATAGTAGC 292

Qy   81 TyrArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGln 100
         |||||||||||||||||||||||||||||||||||||||||||||||||
Db 293 TACAGACAGAGGGCCCGGCTGTTGAAGGACCAGCTCTCCCTGGGAAATGCTGCACTTCAG 352

Qy 101 IleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerTyrGlyGly 120
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 353 ATCACAGATGTGAAATTGCAGGATGCAGGGGTGTACCGCTGCATGATCAGCTATGGTGGT 412

Qy 121 AlaAspTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg 140
         ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 413 GCCGACTACAAGCGAATTACTGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGA 472

Qy 141 IleLeuLeuValAlaAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr 160
         |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 473 ATTTTGCTTGTGGTTGGGGATCCAGTCACCTCTGAACATGAACTGACCTGCCAGGCTGAGGGCTAC 532

Qy 161 ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThrThr 180
         |||||||||||||||||||||||||||||||||||||||||||||||||
Db 533 CCCAAGGCCGAAGTCATCTGGACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCACC 592

Qy 181 ThrThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIleAsn 200
         |||||||||||||||||||||||||||||||||||||||||||||||||
Db 593 ACCACCAATTCCAAGAGAAGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATCAAC 652

Qy 201 ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsnHis 220
         |||||||||||||||||||||||||||||||||||||||||||||||||
Db 653 ACAACAACTAATGAGATTTTCTACTGCACTTTAGGGAGATTAGATCCTGAGGAAAACCAT 712

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Qy  221 ThrAlaGluLeuValIleProGlyLeuProLeuAlaHisProProAsnGluArgThrHis     240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  713 ACAGCTGAATTGGTCATCCCAGAACTACCCTCTGGCACATCCTCCAAATGAAAGGACTCAC   772

Qy  241 LeuValIleLeuGlyAlaIleLeuLeuCysLeuGlyValAlaLeuThrPheIlePheArg     260
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  773 TTGGTAATTCTCGGGGCCATCTTATTATGCCTTGGTGTGACACTGACATTCATCTTCCGT   832

Qy  261 LeuArgLysGlyArgMetMetAspValLysLysCysGlyIleGlnAspThrAsnSerLys     280
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  833 TTAAGAAAAGGGAGAATGATGGATGTGAAAAAATGGCCATCCAAGATACAAACTCAAAG    892

Qy  281 LysGlnSerAspThrHisLeuGluGluThr     290
        ||||||||||||||||||||||||||||||
Db  893 AAGCAAAGTGATACACATTTGGAGGAGACG   922
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RESULT 3
US-09-875-338-1

sequence 1, Application US/09875338
...tent No. US20020055024A1
...NERAL INFORMATION:
APPLICANT: MIKESELL, GLEN E.
APPLICANT: CHANG, HAN
APPLICANT: FINGER, JOSHUA N.
APPLICANT: YANG, GUCHEN
APPLICANT: LU, PIN
APPLICANT: ZHOU, XIA-DI
APPLICANT: PEACH, ROBERT
TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
TITLE OF INVENTION: IMMUNOMODULATION
FILE REFERENCE: 3053-407IUS2
CURRENT APPLICATION NUMBER: US/09/875,338
CURRENT FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/272,107
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: 60/209,811
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 94
SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 1
LENGTH: 1604
TYPE: DNA
ORGANISM: Homo sapiens

Alignment Scores:
Pred. No.:

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...e:                              1.33e-174       Length:          1604
...cent Similarity:               1511.00         Matches:          290
best Local Similarity:            100.00%         Conservative:       0
Query Match:                      100.00%         Mismatches:         0
DB:                               100.00%         Indels:             0
                                  10              Gaps:               0
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US-09-649-108-1 (1-290) x US-09-875-338-1 (1-1604)

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Qy    1 MetArgIlePheAlaValPheIleLeuPheMetThrTyrTrpHisLeuLeuAsnAlaPheThr   20
        ||||||||||||||||||||||||||||||||||||||||||||||||
Db   93 ATGAGGATATTTGCTGTCTTTATATTCATGACCTACTGGCATTTACT    152

Qy   21 ValThrValProLysAspLeuTyrValValGluTyrGlySerAsnMetThrIleGluCys     40
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  153 GTCACGGTTCCCAAGGACCTATATGTGGTAGAGTATGGCAATATGACAATTGAATGC      212

Qy   41 LysPheProValGluLysGlnLeuAspLeuAlaAlaLeuIleValTyrTrpGluMetGlu     60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  213 AAATTCCCAGTAGAAAAACAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAG   272

Qy   61 AspLysAsnIleIleGlnPheValHisGlyGluGluAspLeuLysValGlnHisSerSer     80
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  273 GATAAGAACATTATTCAATTTGTGCATGGAGAAGAAGACCTGAAGGTTCAGCATAGTAGC   332

Qy   81 TyrArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGln    100
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Db  333 TACAGACAGAGGGCCCGGCTGTTGAAGGACCAGCTCTCCCTGGGAAATGCTGCACTTCAG   392

Qy  101 IleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerTyrGlyGly    120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  393 ATCACAGATGTGAAATTGCAGGATGCAGGGGTGTACCGCTGCATGATCAGCTATGGTGGT   452

Qy  121 AlaAspTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg    140
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  453 GCCGACTACAAGCGAATTACTGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGA   512

Qy  141 IleLeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr    160
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  513 ATTTTGGTTGTGGATCCGGTCACCTCTGAACATGAACTGACATGCCAGGCTGAGGGCTAC   572

Qy  161 ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThrThr    180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  573 CCCAAGGCCGAAGTCATCTGGACAAGCAGTGACCATCAAGTGCTGAGTGGTAAGACCACC   632

Qy  181 ThrThrAsnSerLysGluGluLysPheAsnValThrSerThrLeuArgIleAsn        200
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  633 ACAACCAATTCCAAGAGAAGAAGTTTCAATGTGACCAGCACACTGAGAATCAAC        692

Qy  201 ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsnHis    220
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  693 ACAACAACTAATGAAGATTTTTCACTGCACTTTTAGGAGATTAGATCCTGAGGAAAACCAT  752

Qy  221 ThrAlaGluLeuValIleProGlyLeuProLeuAlaHisProProAsnGluArgThrHis    240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  753 ACAGCTGAATTGGTCATCCCAGAACTACCCTCTGGCACATCCTCCAAATGAAAGGACTCAC  812

Qy  241 LeuValIleLeuGlyAlaIleLeuLeuCysLeuGlyValAlaLeuThrPheIlePheArg    260
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  813 TTGGTAATTCTCGGGGCCATCTTATTATGCCTTGGTGTGGCACTGACATTCATCTTCCGT   872

Qy  261 LeuArgLysGlyArgMetMetAspValLysLysCysGlyIleGlnAspThrAsnSerLys    280
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  873 TTAAGAAAAGGGAGAATGATGGATGTGAAAAAATGGCCATCCAAGATACAAACTCAAAG    932

Qy  281 LysGlnSerAspThrHisLeuGluGluThr    290
        |||||||||||||||||||||||||||||||
Db  933 AAGCAAAGTGATACACATTTGGAGGAGACG   962
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RESULT 4
US-09-796-858-41

Sequence 41, Application US/09796858
Patent No. US20020055139A1
GENERAL INFORMATION:
APPLICANT: Holtzmann, Douglas
TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING PROGNOSTIC, DIAGNOSTIC,
TITLE OF INVENTION: PREVENTIVE, THERAPEUTIC, AND OTHER USES
FILE REFERENCE: 7853-226-999
CURRENT APPLICATION NUMBER: US/09/796,858
CURRENT FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 09/223,094
PRIOR FILING DATE: 1998-12-30
PRIOR APPLICATION NUMBER: 09/223,546
PRIOR FILING DATE: 1998-12-30
PRIOR APPLICATION NUMBER: 09/224,246
PRIOR FILING DATE: 1998-12-30
PRIOR APPLICATION NUMBER: 09/312,359
PRIOR FILING DATE: 1999-05-14
PRIOR APPLICATION NUMBER: 09/336,536
PRIOR FILING DATE: 1999-06-18
PRIOR APPLICATION NUMBER: 09/342,687
PRIOR FILING DATE: 1999-06-29
PRIOR APPLICATION NUMBER: 09/399,723
PRIOR FILING DATE: 1999-09-20
PRIOR APPLICATION NUMBER: 09/471,179
PRIOR FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: 09/474,071
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/474,072

PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/572,002
PRIOR FILING DATE: 2000-05-14
PRIOR APPLICATION NUMBER: 09/597,993
PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: 09/599,596
PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 09/606,565
PRIOR FILING DATE: 2000-06-29
PRIOR APPLICATION NUMBER: 09/365,164
PRIOR FILING DATE: 1999-07-30
PRIOR APPLICATION NUMBER: 09/630,334
PRIOR FILING DATE: 2000-07-31
PRIOR APPLICATION NUMBER: 09/665,666
PRIOR FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 50
SEQ ID NO 41
LENGTH: 3575
TYPE: DNA
RGANISM: Homo sapiens
U.

US-09-649-108-1 (1-290) x US-09-796-858-41 (1-3575)

Alignment Scores:
Pred. No.:
Score: 1511.00
Percent Similarity: 100.00%
Percent Identity: 100.00%
Best Local Similarity: 100.00%
Query Match: 10
DB:

Length: 3575
Matches: 290
Conservative: 0
Mismatches: 0
Indels: 0
Gaps: 0

Pred. No.: 4.46e-174

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Qy   1   MetArgIlePheAlaValPheIleGluPheMetThrTyrThrTrpHisLeuLeuAsnAlaPheThr   20
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  59   ATGAGGATATTTGCTGTCTTTATATTCATGACCTACTGGCATTTGCTGAACGGCATTTACT  118

Qy  21   ValThrValProLysAspLeuTyrTyrValValGluTyrTyrGlySerAsnMetThrIleGluCys   40
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 119   GTCACGGTTCCCAAGGACCTATATGTGGTAGAGTATGGTAGCAATATGACAATTGAATGC  178

Qy  41   LysPheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleValTyrThrGlyMetGlu   60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 179   AAATTCCCAGTAGAAAAACAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAG  238

Qy  61   AspLysAsnIleIleGlnPheValHisGlyGluGluAsnGlnHisSerSer   80
         |||||||||||||||||||||||||||||||||||||||||||||||||||
Db 239   GATAAGAACATTATTCAATTTGTGCATGGAGAAGAGAACCAGAAGGTTCAGCATAGTAGC  298

Qy  81   TyrArgGlnAsnGlnAlaArgLeuLeuLysSerGlnLeuSerLeuGlyAsnAlaGlnArg  100
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 299   TACAGACAAGAGGGCCCCGGCTGTTGAAGGACCACCTCTCCCTGGGAAATGCTGCACTTCAG  358

Qy 101   IleThrAspValLysAspAlaGlyValValTyrArgCysMetIleSerTyrGlyGly  120
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 359   ATCACAGATGTGAAAATTGCAGGATGCAGGGGTGTACCGCTGCATGATCATGCTATGGTGGT  418

Qy 121   AlaAspTyrLysArgGlnThrValLysSerValAsnAlaProTyrAsnLysIleLeuAsnGlnArg  140
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 419   GCCGACTACAAGCGACAATACGTGTTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGA  478

Qy 141   IleLeuLeuValAlaAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr  160
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 479   ATTTTGGTTGTGGATCCAGTCACCTCTGAACATGAACTGACATGTCAGGCTGAGGGCTAC  538

Qy 161   ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThrThr  180
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 539   CCCAAGGCCGAAGTCATCTGGACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCACC  598

Qy 181   ThrThrAsnSerLysArgGluGluLysGlnLeuPheAsnValThrSerThrLeuArgLeuAsn  200
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 599   ACACCAATTCCAAGAGAGGAGAAGCAGTTTTCAAATGTGACCAGCACACTGAGAATCAAC  658

Qy 201   ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsnHis  220
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 659   ACAACAACTAATGAGAATTTCTACTGCACTTTTAGGAGATTAGATCCTGAGGAAAACCAT  718

Qy 221   ThrAlaGlyIleLeuValIleLeuProGluProLeuAlaHisProProAsnGluArgThrHis  240
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 719   ACAGCTGGAATTGTCATCCCAGAACTACCTGCTCACCCTCCAAATGAAAGGACTCAC  778

Qy 241   LeuValIleLeuGlyAlaIleLeuLeuCysLeuGlyValAlaLeuThrPheIlePheArg  260
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 779   TTGGTAATTCTGGGAGCCCATCTTATTATGCCTTGGTGTGCTAGCACTGACATTCATCTTCCGT  838

Qy 261   LeuArgLysGlyArgMetMetAspValLysLysCysGlyIleGlnAspThrAsnSerLys  280
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 839   TTAAGAAAAGGGAGAATGATGGACGTGAAAAAATGTGGCATCCAAGATACAAACTCCAAAG  898

Qy 281   LysGlnSerAspThrHisLeuGluGluThr  290
         ||||||||||||||||||||||||||||||
Db 899   AAGCAAAGTGATACACATTGGGAGGAGACG  928
```

RESULT 5
US-09-875-338-3

Sequence 3, Application US/09875338
Patent No. US2002009502A1
GENERAL INFORMATION:
APPLICANT: MIKESELL, GLEN E.
APPLICANT: CHANG, HAN
APPLICANT: FINGER, JOSHUA N.
APPLICANT: YANG, GUCHEN
APPLICANT: LU, PIN
APPLICANT: ZHOU, XIA-DI
APPLICANT: PEACH, ROBERT
TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
TITLE OF INVENTION: IMMUNOMODULATION
FILE REFERENCE: 3053-4071US2
CURRENT APPLICATION NUMBER: US/09/875,338
CURRENT FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/272,107
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: 60/209,811
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 94
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3
LENGTH: 3600
TYPE: DNA
ORGANISM: Homo sapiens
US-09-875-338-3

US-09-649-108-1 (1-290) x US-09-875-338-3 (1-3600)

Alignment Scores:
Pred. No.:
Score: 1511.00
Percent Similarity: 100.00%
Percent Identity: 100.00%
Best Local Similarity: 100.00%
Query Match: 10
DB:

Length: 3600
Matches: 290
Conservative: 0
Mismatches: 0
Indels: 0
Gaps: 0

Pred. No.: 4.51e-174

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Qy  1   MetArgIlePheAlaValPheIleGluPheMetThrTyrThrTrpHisLeuLeuAsnAlaPheThr  20
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 93   ATGAGGATATTTGCTGTCTTTATATTCATGACCTACTGGCATTTGCTGAACGGCATTTACT  152

Qy 21   ValThrValProLysAspLeuTyrTyrValValGluTyrTyrGlySerAsnMetThrIleGluCys  40
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 153  GTCACGGTTCCCAAGGACCTATATGTGGTAGAGTATGGTAGCAATATGACAATTGAATGC  212

Qy 41   LysPheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleValTyrThrGlyMetGlu  60
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 213  AAATTCCCAGTAGAAAAACAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAG  272

Qy 61   AspLysAsnIleIleGlnPheValHisGlyGluGluAsnGlnHisSerSer  80
        ||||||||||||||||||||||||||||||||||||||||||||||||||
Db 273  GATAAGAACATTATTCAATTTGTGCATGGAGAAGAGAACCAGAAGGTTCAGCATAGTAGC  332
```

```
Qy      81 TyrArgGlnArgAlaArgLeuLeuLysSerLeuSerLeuGlyAsnAlaAlaLeuGln 100
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     333 TACAGACAGAGGGCGCGGCTGTTGAAGAGCCAGCTCTCCCTGGGAAATGCTGCACTTCAG 392

Qy     101 IleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerThrTyrGlyGly 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     393 ATCACAGATGTGAAATTGCAGGATGCAGGGGTGTACCGGTGCATGATCAGCTATGGTGGT 452

Qy     121 AlaAspThrTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg 140
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     453 GCCGACTACAAGAGGATTACTGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGA 512

Qy     141 IleLeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr 160
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     513 ATTTTGGTTGTGGATCCAGTCACCTCTGAACATGAACTGACATGTCAGGCTGAGGGCTAC 572

Qy     161 ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThrThr 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     573 CCCAAGGCCGAAGTCATCTGGACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCACC 632

Qy     181 ThrThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIleAsn 200
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     633 ACCACCAATTCCAAGAGAGAAGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATCAAC 692

Qy     201 ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsnHis 220
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     693 ACAACAACTAATGAGATTTTCTACTGCACTTTTAGGAGATTAGACCCTGAGGAAAACCAT 752

Qy     221 ThrAlaGluGluValIleProGlyProLeuAlaHisProProAsnGluArgThrHis 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     753 ACAGCTGAATTGGTCATCCCAGAACTACCCTCGCCACATCCTCCAAATGAAAGGACTCAC 812

Qy     241 LeuValIleLeuGlyValAlaIleLeuLeuCysValAlaLeuThrPheIlePheArg 260
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     813 TTGGTAATTCTGGGAGCCATCTTATTATGTGTAGCACTGACATTCATCTTCCGT 872

Qy     261 LeuArgLysGlyArgMetMetAspValLysLysCysGlyIleGlnAspThrAsnSerLys 280
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     873 TTAAGAAAAGGGAGAATGATGGATGTGAAAAAATGTGGCATCCAAGATACAAACTCAAG 932

Qy     281 LysGlnSerAspThrHisLeuGluThr 290
           ||||||||||||||||||||||||||
Db     933 AAGCAAAGTGATACACATTTGGAGGAGACG 962
```

RESULT 6
US-10-002-775-1
Sequence 1, Application US/10002775
Patent No. US20020102651A1
GENERAL INFORMATION:

APPLICANT: Gordon Freeman
APPLICANT: Vassiliki Boussiotis
APPLICANT: Tatyana Chernova
APPLICANT: Nelly Malenkovich
TITLE OF INVENTION: NOVEL B7-4 MOLECULES AND USES THEREFOR
FILE REFERENCE: GNN-004ADV
CURRENT APPLICATION NUMBER: US/10/002,775
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: US 09/644,934
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 60/150,390
PRIOR FILING DATE: 1999-08-23
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentln Ver. 2.0

SEQ ID NO 1
LENGTH: 968
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (59)..(793)
US-10-002-775-1

Alignment Scores:
Pred. No.:
Score: 4.69e-135 Length: 968
Percent Similarity: 1184.00 Matches: 227
Best Local Similarity: 100.00% Conservative: 0
Query Match: 100.00% Mismatches: 0
DB: 78.36% Indels: 0
                           12 Gaps: 0

US-09-649-108-1 (1-290) x US-10-002-775-1 (1-968)

```
Qy       1 MetArgIlePheAlaValPhoIlePheMetThrTyrTrpHisLeuLeuAsnAlaPheThr 20
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      59 ATGAGGATATTTGCTGTCTTTATATTCATGACCTACTGGCATTTGCTGAACGCATTTACT 118

Qy      21 ValThrValProLysAspLeuTyrValValGluTyrGlySerAsnMetThrIleGluCys 40
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     119 GTCACGGTTCCCAAGGACCTATATGTGGTAGAGTATGGTAGCAATATGACAATTGAATGC 178

Qy      41 LysPheProValGlnLysLysGlnLeuAspLeuAlaAlaLeuIleValAlaTyrTrpGluMetGlu 60
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     179 AAATTCCCAGTAGAAAAAAAACAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAG 238

Qy      61 AspLysAsnIleIleGlnPheValHisGlyGluGluAspLeuLysValGlnHisSerSer 80
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     239 GATAAGAACATTATTCAATTCGTGCATGGAGAGGAAGACCTGAAGGTTCAGCATAGTAGC 298

Qy      81 TyrArgGlnArgAlaArgLeuLeuLysSerLeuSerLeuGlyAsnAlaAlaLeuGln 100
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     299 TACAGACAGAGGGCGCGGCTGTTGAAGAGCCAGCTCTCCCTGGGAAATGCTGCACTTCAG 358

Qy     101 IleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerThrTyrGlyGly 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     359 ATCACAGATGTGAAATTGCAGGATGCAGGGGTGTACCGGTGCATGATCAGCTATGGTGGT 418

Qy     121 AlaAspThrTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg 140
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     419 GCCGACTACAAGAGGATTACTGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGA 478

Qy     141 IleLeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr 160
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     479 ATTTTGGTTGTGGATCCAGTCACCTCTGAACATGAACTGACATGTCAGGCTGAGGGCTAC 538

Qy     161 ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThrThr 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     539 CCCAAGGCCGAAGTCATCTGGACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCACC 598

Qy     181 ThrThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIleAsn 200
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     599 ACCACCAATTCCAAGAGAGAAGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATCAAC 658

Qy     201 ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsnHis 220
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     659 ACAACAACTAATGAGATTTTCTACTGCACTTTTAGGAGATTAGACCCTGAGGAAAACCAT 718

Qy     221 ThrAlaGluGluValIlePro 227
           |||||||||||||||||||||
Db     719 ACAGCTGAATTGGTCATCCCA 739
```

RESULT 7
US-09-875-338-4
Sequence 4, Application US/09875338
Patent No. US20020095024A1
GENERAL INFORMATION:

APPLICANT: MIKESELL, GLEN E.
APPLICANT: CHANG, HAN
APPLICANT: FINGER, JOSHUA N.
APPLICANT: YANG, GUCHEN
APPLICANT: LU, PIN
APPLICANT: ZHOU, XIA-DI
APPLICANT: PEACH, ROBERT
TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
TITLE OF INVENTION: IMMUNOMODULATION
FILE REFERENCE: 3053-4071US2

CURRENT APPLICATION NUMBER: US/09/875,338
CURRENT FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/272,107
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: 60/209,811
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 94
SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 4
LENGTH: 1443
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: fusion construct
US-09-875-338-4

Alignment Scores:
Pred. No.:   1.07e-128   Length: 1443
S...:   1114.00   Matches: 217
...nt Similarity: 100.00%   Conservative: 1
Best Local Similarity: 99.548%   Mismatches: 0
Query Match: 10   Indels: 0
                                       Gaps: 0

US-09-649-108-1 (1-290) x US-09-875-338-4 (1-1443)

```
Qy  22 ThrValProLysAspLeuTyrValValGluTyrGlySerAsnMetThrIleGluCysLys  41
       |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  76 AGTGTTCCCAAGGACCTATATGTGGTAGAGTATGGTAGCAATATGACAAATTGAAATGCAAA 135

Qy  42 PheProValGlyLeuValSerLeuAsnLeuAlaLeuLeuValLeuValThrTyrTrpGluMetGluAsp 61
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 136 TTCCCAGTAGGAAAAACAAATTAGACCTGGCTGCACTAATTGTCTATTGGGAAATGGAGGAT 195

Qy  62 LysAsnIleIleGlnHisPheValHisGlyGluGluAspSerTyr                 81
       |||||||||||||||||||||||||||||||||||||||||||||
Db 196 AAGAACATTATTCAATTTGTGCATGGAGAAGAAGACTTCAGCTAC                255

Qy  82 ArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGlnIle 101
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 256 AGACAGAGGGCCCGGCTGTTGAAGGACCAGCTCTCCCTGGGAAATGCTGCACTGCAGATC 315

Qy 102 ThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerTyrGlyGlyAla 121
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 316 ACAGATGTGAAATTGCAGGATGCAGGGGTGTACCGGTGCATGATCAGCTATGGTGGTGCC 375

Qy 122 AspTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArgIle 141
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 376 GACTACAAGCGAATTACTGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGAATT 435

Qy 142 LeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyrPro 161
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 436 TTGGTTGTGGATCCAGTCACCTCTGAACATGAACTGACATGTCAGGCTGAGGGCTACCCC 495

Qy 162 LysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyGlyLysThrThr 181
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 496 AAGGCCGAAGTCATCTGGACAAGCAGCGACCATCAAGTCCTGAGTGGTGGAAGACCACC   555

Qy 182 ThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIleAsnThr 201
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 556 ACCAATTCCAAGAGAGAGGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATCAACACA 615

Qy 202 ThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluAsnHisThr     221
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 616 ACAACTAATGAGATTTTCTACTGCACTTTAGGGAGATTAGATCCTGAGGAAAACCATACA 675

Qy 222 AlaGluLeuValIleProGluProLeuAlaHisProProAsnGluArgThr          239
       |||||||||||||||||||||||||||||||||||||||||||
Db 676 GCTGAATTGGTCATCCCAGAACTACCTCTCCAAATGAAAGGACT                729
```

Sequence 10, Application US/10002775
Patent No. US20020102651A1
GENERAL INFORMATION:
APPLICANT: Gordon Freeman
APPLICANT: Vassiliki Boussiotis
APPLICANT: Tatyana Chernova
APPLICANT: Nelly Malenkovich
TITLE OF INVENTION: NOVEL B7-4 MOLECULES AND USES THEREFOR
FILE REFERENCE: GNN-004ADV
CURRENT APPLICATION NUMBER: US/10/002,775
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: US 09/644,934
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 60/150,390
PRIOR FILING DATE: 1999-08-23
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 10
LENGTH: 3593
TYPE: DNA
ORGANISM: Mus musculus
FEATURE:
NAME/KEY: CDS
LOCATION: (17)..(889)
US-10-002-775-10

Alignment Scores:
Pred. No.:   7.42e-118   Length: 3593
Score: 1050.00   Matches: 202
Percent Similarity: 81.10%   Conservative: 34
Best Local Similarity: 69.42%   Mismatches: 53
Query Match: 12   Indels: 2
                                       Gaps: 2

US-09-649-108-1 (1-290) x US-10-002-775-10 (1-3593)

```
Qy   1 MetArgIlePheAlaValPheIlePheMetThrTyrTrpHisLeuLeuAsnAlaPheThr  20
       ||||||||||||||  |||||| :::: || ::::::::||||  ||||||||||||
Db  17 ATGAGGATATTTGCTGGCATTATATTCACAGCCTGCTGCCTACTGGCCGTTTACT      76

Qy  21 ValThrValProLeuTyrValValGlyTyrGlySerAsnMetThrIleGluCys       40
       ||:||:|  ||||||||||||||||||||||||||||||||:::||::|||||||
Db  77 ATCACGGCCTCCAAAGGACTTGTACCTGGTGGAGTATGGCAGCAACGTCACGATGGAGTGC 136

Qy  41 LysPheProValGlyLeuGlnAspAlaGlyValTyrArgCysMetIleSerTyrGlyGly  60
       |||||||||||||  :::::||||||||||||||||||||||||||||||||||||||
Db 137 AGATTCCCTGTAGGAACGGAGACCTGCTTGTGGTGTACTGGGAAAAAGGAA          196

Qy  61 AspLysAsnIleIleGlnHisPheValHisGlyGluGluAspSerLeuHisSerSer     80
       |||:::|||||||||||||||||||||||||||||||||||||:::||::
Db 197 GATGAGCAAGTGATTCAGTTTGTGGCAGGAGGAGGACCTTAAGCCTACAGCACAGCCAAC 256

Qy  81 TyrArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGln 100
       ::||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 257 TTCAGGAGGAGAGCCTCGCTGCCAAAGGACCAGCTTTTGAAGGAAATGTGCCCCTTCAG  316

Qy 101 IleThrAspValLysLeuGlnAspAlaGlyValTyrArgCysMetIleSerTyrGlyGly 120
       |||||||||:|:::||||||||||||||||||||||||||||||||||||||||||||||
Db 317 ATCACAGACGTCAAGCTGCAGGACGCCAGGGCGTTTACTGTGCCATAATCAGCTACGGTGGT 376

Qy 121 AlaAspTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg 140
       ||||||||||||||:||||||::||||:||||||||||||||||||||||||||||||||
Db 377 GCGGACTACAAGCGAATCACCGTCAGAAAGTCAATGCCCCATACAACAAAATCAACCAGAGA 436

Qy 141 IleLeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr 160
       |||   |:||||||||||||||||||||||||::||||||||||||||||||||||||||
Db 437 ATT---TCCGTGGATCCAGCCACTTCTGAGCAATGAACTAATATGTCAGGCCGAGGGTTAT 493

Qy 161 ProLysAlaGluValIleTrpThrSerSerAspHisGlnValLeuSerGlyGlyLysThr 180
       ||||||||||||||||||||||||::||||||||||||:::|||||||||||||||||||
Db 494 CCAGAAGCTGAGGTAATCTGGACAAACAGTGACCACCAACCGTGAGTGGGGAAGACAAGT  553

Qy 181 ThrThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIleAsn 200
       ||||||||||||||||||||||||||||||||||||
Db 181 ...                                                         200
```

RESULT 8
US-10-002-775-10

```
Db  554 GTCACCACTTCCCGacCAGAGGGGGATGCTTCTCAANGTGACCAGCAGTCTGAGGGTCAAC 613
            |||   |||::::   |||                |||||||||||::::||
Qy  201 ThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGlnAsnHis 220

Db  614 GCCACAGCGAAATGATGTTTTCTACGTTACGTTTGGGAATCACGCCAGGGCAAAACCAC 673
            |||   ::||||||||||||||||||||||||||           :||||||
Qy  221 ThrAlaGlyGluLeuValIleProGluLeuProLeuAlaHisProProAsnGluArgThrHis 240

Db  674 ACACGGAGCTGATCATCCCAGAACTGCCTGCAACACATCCTTCCACAGAACAGGACTCAC 733
            ||||||||||          ::||||||||||||
Qy  241 LeuValIleLeuGlyAlaIleAlaIleLeuLeuCysLeuGlyValAlaAlaLeuThrPheArg 260

Db  734 TGGGTGCTTCTGGGATCCATCCTGTTGTTCCCATTGTAGTGTCCACGGGTCCTCCTCTTC 793
            |||||||||||||||||||||         :||||||
Qy  261 LeuArgLysGly--ArgMetMetAspValLysLysCysGlyIleGlnAspThrThrAsnSer 279

Db  794 TTGAGAAAACAAGTGAGAATGCTAGAGTGTCGGAGAAATGTGGCGTTGAAGATACAAGCTCA 853
            |||||||||||||||::::|||||::::||||||||||::::||
Qy  280 LysLysGlnSerAspThrHisLeuGluGlnThr 290

Db  854 AAAAACCGAAATGATACACAAATTCGAGGAGACG 886
            :::||||||||
```

RESULT 9

US-09-796-858-43

Sequence 43, Application US/09796858

Patent No. US2002055139A1

GENERAL INFORMATION:
APPLICANT: Holtzmann, Douglas
TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING PROGNOSTIC, DIAGNOSTIC,
TITLE OF INVENTION: PREVENTIVE, THERAPEUTIC, AND OTHER USES
FILE REFERENCE: 7853-226-999
CURRENT APPLICATION NUMBER: US/09/796,858
CURRENT FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 2001-03-01
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: 09/665,666
PRIOR FILING DATE: 2000-07-31
PRIOR APPLICATION NUMBER: 09/630,334
PRIOR FILING DATE: 1999-07-30
PRIOR APPLICATION NUMBER: 09/365,164
PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 09/606,565
PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: 09/599,596
PRIOR FILING DATE: 2000-05-14
PRIOR APPLICATION NUMBER: 09/597,993
PRIOR FILING DATE: 2000-06-29
PRIOR APPLICATION NUMBER: 09/572,002
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/474,072
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/474,071
PRIOR FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: 09/471,179
PRIOR FILING DATE: 1999-09-20
PRIOR APPLICATION NUMBER: 09/299,723
PRIOR FILING DATE: 1999-06-29
PRIOR APPLICATION NUMBER: 09/342,687
PRIOR FILING DATE: 1999-06-18
PRIOR APPLICATION NUMBER: 09/336,536
PRIOR FILING DATE: 1999-05-14
PRIOR APPLICATION NUMBER: 09/312,359
PRIOR FILING DATE: 1998-12-30
PRIOR APPLICATION NUMBER: 09/224,246
PRIOR FILING DATE: 1998-12-30
PRIOR APPLICATION NUMBER: 09/223,094
PRIOR FILING DATE: 1998-12-30
PRIOR APPLICATION NUMBER: 09/223,546

FEATURE:
NAME/KEY: modified_base
LOCATION: all "n" positions
OTHER INFORMATION: n=a, c, g, or t
US-09-796-858-43

Alignment Scores:

Pred. No.;	1.38e-103	Length: 891
Score:	925.50	Matches: 191
Percent Similarity:	79.36%	Conservative: 32
Best Local Similarity:	67.97%	Mismatches: 54
Query Match:	61.25%	Indels: 7
DB:	10	Gaps: 3

US-09-649-108-1 (1-290) x US-09-796-858-43 (1-891)

```
Qy    1 MetArgIlePheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleLeuValTyrTrpGluMetGlu 20
          ||||||||||||||||||||||||||||||||||||
Db    1 AGATTCCCTGTAGAACGGGAAGCTGGACCTGCTGTTGCGTTAGTGTGTTACTGGGAAAAGGAA 234

Qy   21 ValThrValProLysAspGluLeuTyrValValGluGlyTyrGlySerAsnMetThrIleGluCys 40
          :::|||       |||||||||||||||||||||::::|||::::|||
Db   55 ATGGAGATATTTTGCTGGCCATTATATTCACAGCCTGCTGTCACTTGCTACGGGCGGTTACT 114

Qy   41 LysPheProValGlyLysGlnLeuAspLeuAlaAlaLeuIleLeuValTyrTrpGluMetGlu 60
          |||||||||||||||
Db  115 ATCACGGCTCCAAAGGACTTGTACCGTGGTGGAGTATGGCAGAGAACGTCACGATGGAGTGC 174

Qy   61 AspLysAsnIleIleGlnPheValHisGlyGluGluAspLeuLysValAspAsnHisSer 80
          |||:::|||       |||||||
Db  175 ATGGAGATATTTTGCTGGCCATTATATTCACAGCCTGCTGTCACTTGCTACGGGCGGTTACT 234

Qy   81 TyrArgGlnArgAlaArgLeuLeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGln 100
          ||||||||||||||||||
Db  235 GATGAGCAAGTGATCATCAGTTTGTGGCAGAGGAGGAGGACCTTAAGCCTCAGACAGCAAC 294

Qy  101 IleThrAspValLysLysLeuGluIleAspAsnAlaGlyValTyrArgCysMetIleSerTyrGlyGly 120
          ||::::|||||||
Db  295 TTCAGGGGGAGAGCCTCGCTTGCCAAAGGACCCAGCTTTGAAGGAAATGCTGCCTTCAG 354

Qy  121 AlaAspTyrLysLeuArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGlnArg 140
          ||||||||||||||||||::::|||
Db  355 ATCACAGACGTCAAGCTGCCAGACGCGTTACTGCTGCATATAATCAGCTACGGTGGT 414

Qy  141 IleLeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGlyTyr 160
          ||||||::::|||||||
Db  415 GCGGACTACAAGCGAATCACGGTGAAGTCAATGCCCCATACCGCAAAATCAACCAGAGA 474

Qy  161 ProLysSerAlaGluLeuThrPheThrSerSerAspHisGlnValLeuSerGlyLysThrThr 180
          ||::::|||||||::::|||
Db  475 ATT---TCCGTGGATCCAGCCACTTCTGAGCATGAACTAATATGTCAGGCCGAGGGTTAT 531

Qy  181 ThrThrAsnSerLysArgGluLysLysGlnSerAspThrHisSerThrArgLeuArgIleAsn 200
          |||::::::
Db  532 CCAGAAGCTGAGTAATCTAGCAAAGTGACCACCAACCCGTAGTGTGGGAAGAGAAGT 591

Qy  201 ThrThr------ThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGlu 218
          |||::::
Db  592 GTCACCACTTCCCGGACAGAGGGGGATGCTTCTCAAATGACCAGCAGTCTC-AAC 650

Qy  219 AsnHisThrAlaGluLeuValIleProGluLeuProLeuAlaHisProProAsnGluArg 238
          ||||||:::::
Db  651 GCCACAGCGAANNAGCGAATGATGTT-TTCTACGTTACGTTTGGGAATCACGCCAGGGCAA 709

Qy  239 ThrHisLeuValIleLeuGlyAlaIleAlaIleLeuLeuCysLeuGlyValAlaAlaLeuThrPheIle 258
          |||:::|||::::
Db  710 AACCACACAGCCGGC-GANATCACCCCAGAACTGCCTGCAACACATCCTTCCACAGAACAGG 768

Qy  259 PheArgLeuArgLysGly---ArgMetMetAspValLysLysCysGlyIleGlnAspThr 277
          ||||||||||::::||
Db  769 ACTCACTGGGGTGCTTCTGGGATCCATCCTGTTGTTCCCATTGTAGTGTCCACGGGTCCTC 828

Qy  278 Asn 278
          |||
Db  829 CTCTTCTTGAGAAAACAAGTGAGAAATCTAGAGTGTGAGAAAATGTGGCGTTGAAGATACA 888
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NUMBER OF SEQ ID NOS: 50
SEQ ID NO 43
LENGTH: 891
TYPE: DNA
ORGANISM: Mus musculus

Db   889 AGC 891

:::

RESULT 10
US-09-867-701-10728
Sequence 10728, Application US/0987701
Patent No. US20020132237A1
GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10728
ENGTH: 464
YPE: DNA
ORGANISM: Homo sapien
US-09-867-701-10728

Alignment Scores:
Pred. No.:   1.1e-79
Score: 726.00   Length: 464
Percent Similarity: 100.00%   Matches: 140
Best Local Similarity: 100.00%   Conservative: 0
Query Match: 48.05%   Mismatches: 0
DB: 10   Indels: 0
                                      Gaps: 0

US-09-649-108-1 (1-290) x US-09-867-701-10728 (1-464).

Qy   88 LeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGlnIleThrAspValLysLeuGln 107
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   9 TTGAAGGACCAGCTCTCCCTGGGAAATGCTGCACTTCAGATCACAGATGTGAAATTGCAG 68

Qy   108 AspAlaGlyValTyrArgCysMetIleSerTyrGlyGlyAlaAspTyrLysArgIleTh 127
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   69 GATGCAGGGGTGTACCGCTGCATGATCAGCTATGGTGGTGCCGACTACAAGCGAATTACT 128

Qy   128 ValLysValAsnAlaProTyrAsnLysIleAsnGlnArgIleLeuValValAspProVal 147
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   129 GTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGAATTTTGGTTGTGGATCCAGTC 188

Qy   148 ThrSerGluHisGlnLeuThrCysGlnAlaGlnGlyTyrProLysAlaGluValIleTrp 167
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   189 ACCTCTGAACATGAACCTGACATGTCAGGCTGCACCCAAGGCCGAAGTCATCTGG 248

Qy   168 ThrSerSerAspHisGlnValLeuSerGlyLysThrThrThrThrAsnSerLysArgGlu 187
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   249 ACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCACCACCACCAATTCCAAGAGAGAG 308

Qy   188 GluLysLeuPheAsnValThrSerThrLeuArgIleAsnThrThrAsnGluIlePhe 207
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   309 GAGAAGCTTTTCAATGTGACCAGCACTCTGAGAATCAACACAACTAATGAGATTTTC 368

Qy   208 TyrCysThrPheArgArgLeuAspProGluGluAsnHisThrAlaGluLeuValIlePro 227
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   369 TACTGCACTTTTAGGAGATTAGATCCTGAGGAAAACCATACACACTGAATTGGTCATCCCA 428

RESULT 11
US-09-867-701-2957
Sequence 2957, Application US/0987701
Patent No. US20020132237A1
GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2957
LENGTH: 497
TYPE: DNA
ORGANISM: Homo sapien
US-09-867-701-2957

Alignment Scores:
Pred. No.:   4.69e-78
Score: 713.00   Length: 497
Percent Similarity: 99.29%   Matches: 140
Best Local Similarity: 99.29%   Conservative: 0
Query Match: 47.19%   Mismatches: 0
                                      Indels: 1
                                      Gaps: 0

US-09-649-108-1 (1-290) x US-09-867-701-2957 (1-497)

Qy   88 LeuLysAspGlnLeuSerLeuGlyAsnAlaAlaLeuGlnIleThrAspValLysLeuGln 107
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   9 TTGAAGGACCAGCTCTCCCTGGGAAATGCTGCACTTCAGATCACAGATGTGAAATTGCAG 68

Qy   108 AspAla-GlyValTyrArgCysMetIleSerTyrGlyGlyAlaAspTyrLysArgIleTh 127
          |||||| |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   69 GATGCAGGGGTGTACCGCTGCATGATCAGCTATGGTGGTGCCGACTACAAGCGAATTAC 128

Qy   127 rValLysValAsnAlaProTyrAsnLysIleAsnGlnArgIleLeuValValAspProVa 147
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   129 TGTGAAAGTCAATGCCCCATACAACAAAATCAACCAAAGAATTTTGGTTGTGGATCCAGT 188

Qy   147 lThrSerGluHisGlnLeuThrCysGlnAlaGlnGlyTyrProLysAlaGluValIleTr 167
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   189 CACCTCTGAACATGAACCTGACATGTCAGGCTGCACCCAAGGCCGAAGTCATCTG 248

Qy   167 pThrSerSerAspHisGlnValLeuSerGlyLysThrThrThrThrAsnSerLysArgGl 187
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   249 GACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACCACCACCACCAATTCCAAGAGAGA 308

Qy   187 uGluLysLeuPheAsnValThrSerThrLeuArgIleAsnThrThrThrAsnGluIlePh 207
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   309 GGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATCAACACAACTAATGAGATTTT 368

Qy   207 eTyrCysThrPheArgArgLeuAspProGluGluAsnHisThrAlaGluLeuValIlePr 227
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   369 CTACTGCACTTTTAGGAGATTAGATCCTGAGGAAAACCATACAGCGTGAATTGGTCATCCC 428

Qy   227 o 227
          |
Db   429 A 429

RESULT 12
US-09-867-701-3638/c
Sequence 3638, Application US/0987701
Patent No. US20020132237A1
GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3638
LENGTH: 442
TYPE: DNA
ORGANISM: Homo sapien
US-09-867-701-3638

Alignment Scores:
Pred. No.:
Score: 565.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 37.39%

```
Length:        442
Matches:       108
Conservative:  0
Mismatches:    0
Indels:        0
Gaps:          0
```

US-09-649-108-1 (1-290) x US-09-867-701-3638 (1-442)

```
Qy  120 GlyAlaAspThrTyrLysArgIleThrValLysValAsnAlaProTyrAsnLysIleAsnGln 139
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  442 GGTGCCGACTACAAGCGAATTACTGTGAAAGTCAATGCCCAATACAACAAAATCAACCAA 383

Qy  140 ArgIleLeuValValAspProValThrSerGluHisGluLeuThrCysGlnAlaGluGly 159
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  382 AGAATTTTGGTTGTGGATCCAGTCACCTCTGACATGAACTGACATGTCAGGCTGAGGGC 323

Qy  160 TyrProLysAlaGlyValIleIleTrpThrSerSerAspHisGlnValLeuSerGlyLysThr 179
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  322 TACCCCAAGGCCGGAAGTCATCTGGACAAGCAGTGACCATCAAGTCCTGAGTGGTAAGACC 263

Qy  180 ThrThrThrAsnSerLysArgGluGluLysLeuPheAsnValThrSerThrLeuArgIle 199
            |||||||||||||||||||||||||||||||||||||||||||||||||
Db  262 ACCACCACCAATTCCAAGAGAGAGAAGCTTTTCAATGTGACCAGCACACTGAGAATC 203

Qy  200 AsnThrThrThrAsnGluIlePheTyrCysThrPheArgArgLeuAspProGluGluAsn 219
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  202 AACACAACCAACTAATGACGATTTTCTACTCGCACTTTAGGAGATTAGATCCTGAGGAAAAC 143

Qy  220 HisThrAlaGluLeuValIlePro 227
            ||||||||||||||||||||||||
Db  142 CATACACAGCTGAATTGGTCATCCCA 119
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RESULT 13

US-09-875-338-18

; Patent No. US20020095024A1
; Sequence 18, Application US/0975338
; GENERAL INFORMATION:
;   APPLICANT: MIKESELL, GLEN E.
;   APPLICANT: CHANG, HAN
;   APPLICANT: FINGER, JOSHUA N.
;   APPLICANT: YANG, GUCHEN
;   APPLICANT: LU, PIN
;   APPLICANT: ZHOU, XIA-DI
;   APPLICANT: PEACH, ROBERT
;   TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
;   TITLE OF INVENTION: IMMUNOMODULATION
;   FILE REFERENCE: 3053-4071US2
;   CURRENT APPLICATION NUMBER: US/09/875,338
;   CURRENT FILING DATE: 2001-06-06
;   PRIOR APPLICATION NUMBER: 60/272,107
;   PRIOR FILING DATE: 2001-02-28
;   PRIOR APPLICATION NUMBER: 60/209,811
;   PRIOR FILING DATE: 2000-06-06
;   NUMBER OF SEQ ID NOS: 94
;   SOFTWARE: Patentin Ver. 2.1
;   SEQ ID NO 18
;   LENGTH: 666
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-875-338-18

Alignment Scores:
Pred. No.:
Score: 527.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 34.88%

```
Length:        666
Matches:       101
Conservative:  0
Mismatches:    0
Indels:        0
Gaps:          0
```

US-09-649-108-1 (1-290) x US-09-875-338-18 (1-666)

```
Qy  190 LeuPheAsnValThrSerThrLeuArgIleAsnThrThrThrAsnGluIlePheTyrCys 209
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    3 CTTTTCAATGTGACCAGCACACTGAGAATCAACACAACCAACTAATGAGATTTTCTACTGC 62

Qy  210 ThrPheArgArgLeuAspProGluGluAsnHisThrAlaGluLeuValIleProGluLeu 229
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   63 ACTTTTAGGAGATTAGATCCTGAGGAAAACCATACACAGCTGAATTGGTCATCCCAGAACTA 122

Qy  230 ProLeuAlaHisProProAsnGluArgThrHisLeuValIleLeuGlyAlaIleLeuLeu 249
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  123 CCTCTGGCACATCCTCCAAATGAAAGGACTCACTTCGTAATTCTGGGAGCCATCTTATTA 182

Qy  250 CysLeuGlyValAlaLeuThrPheIlePheArgLeuArgLysGlyArgMetMetAspVal 269
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  183 TGCCTTGGTGTGGCTCTGACCTTCATTTTCCGTTTAAGAAAAGGGAGAAATGGGATGGATGTG 242

Qy  270 LysLysCysGlyIleGlnAsnThrAsnSerLysLysGlnSerAspThrHisLeuGluGlu 289
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  243 AAAAAATGGGCATCCAAGATACAAAACTCAAAGAAGCAAAGTGATACACATTTGGAGGAG 302

Qy  290 Thr 290
            |||
Db  303 ACG 305
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RESULT 14

US-09-896-913A-3

; Patent No. US20020164600A1
; Sequence 3, Application US/0896913A
; GENERAL INFORMATION:
;   APPLICANT: Freeman, Gordon
;   APPLICANT: Chernova, Irene
;   APPLICANT: Chernova, Tatyana
;   APPLICANT: Malenkovich, Nelly
;   APPLICANT: Wood, Clive
;   TITLE OF INVENTION: PD-L2 MOLECULES: NOVEL PD-1 LIGANDS AND
;   TITLE OF INVENTION: USES THEREFOR
;   FILE REFERENCE: GNN-026A
;   CURRENT APPLICATION NUMBER: US/09/896,913A
;   CURRENT FILING DATE: 2002-04-15
;   PRIOR APPLICATION NUMBER: 60/214,563
;   PRIOR FILING DATE: 2000-06-28
;   PRIOR APPLICATION NUMBER: 60/270,822
;   PRIOR FILING DATE: 2001-02-23
;   PRIOR APPLICATION NUMBER: 60/271,114
;   PRIOR FILING DATE: 2001-02-23
;   NUMBER OF SEQ ID NOS: 12
;   SOFTWARE: FastSEQ for Windows Version 4.0
;   SEQ ID NO 3
;   LENGTH: 819
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)...(819)
US-09-896-913A-3

Alignment Scores:
Pred. No.:
Score: 411.50
Percent Similarity: 53.14%
Best Local Similarity: 39.11%
Query Match: 27.23%

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Length:        819
Matches:       106
Conservative:  38
Mismatches:    93
Indels:        34
Gaps:          8
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US-09-649-108-1 (1-290) x US-09-896-913A-3 (1-819)

```
Qy   39 GluCysLysPheProValGluLysGlnLeuAspLeuAlaAlaLeuIleValThrTyrTrpGlu 58
            | ||||| |||       ||| |  ::||  |||::
Db  121 GAATGCAACTTTGACACTGGAAGTCAATGTGAAGAAGCCAGTTGCAA 180
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Alignment Scores:

Pred. No.:	
Score:	411.50
Percent Similarity:	53.14%
Best Local Similarity:	39.11%
Query Match:	27.23%
DB:	

Length:	819
Matches:	106
Conservative:	38
Mismatches:	93
Indels:	34
Gaps:	8

Length: 5.82e-41

US-09-649-108-1 (1-290) x US-09-794-210-1 (1-819)

[This page contains a nucleotide/protein sequence alignment between query (Qy) US-09-649-108-1 and database (Db) US-09-794-210-1, presented in two mirrored copies across the page. The alignment consists of paired Query and Database lines with conservation bars (|||, :::) showing matched amino acid and DNA sequences from MetGluAspLys... through the full sequence length.]

Search completed: January 12, 2003, 12:44:57
Job time : 68 secs

OM protein - protein search, using sw model

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GenCore version 5.1.3

Run on: January 12, 2003, 10:10:43 ; Search time 10.5287 Seconds
(without alignments)
480.942 Million cell updates/sec

Title: US-09-649-108-1_copy_30_290
Perfect score: 1356
Sequence: 1 VEYGSNMTIECKFPVEKQLD..........KCGIQDTNSKKQSDTHLEET 261

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 118974 seqs, 19401057 residues

(1) number of hits satisfying chosen parameters: 118974
Listing first 45 summaries

Post-processing: Minimum Match 0%
Maximum Match 100%
Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Database :   Published_Applications_AA.*

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 2:  /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
 3:  /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
 4:  /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
 5:  /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB_NEW_PUB.pep:*
 6:  /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
 7:  /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
 8:  /cgn2_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*
 9:  /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
10:  /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
11:  /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
12:  /cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
13:  /cgn2_6/ptodata/2/pubpaa/US60_NEW_PUB.pep:*
14:  /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1356	100.0	290	9	US-10-068-215-4	Sequence 4, Appli
2	1356	100.0	290	9	US-09-896-738-12	Sequence 12, Appl
3	1356	100.0	290	9	US-09-896-913A-12	Sequence 12, Appl
4	1356	100.0	290	9	US-09-915-789A-17	Sequence 17, Appl
5	1356	100.0	290	10	US-09-796-858-42	Sequence 42, Appli
6	1356	100.0	290	10	US-09-875-338-2	Sequence 2, Appli
7	1356	100.0	290	10	US-09-910-174A-8	Sequence 8, Appli
8	1356	100.0	290	10	US-09-955-866-6	Sequence 6, Appli
9	1356	100.0	290	10	US-09-899-837-12	Sequence 12, Appl
10	1356	100.0	290	12	US-10-002-775-4	Sequence 4, Appl
11	1104	81.4	220	9	US-09-915-789A-23	Sequence 23, Appl
12	1096	80.8	480	10	US-09-875-338-5	Sequence 5, Appli
13	1029	75.9	245	12	US-10-002-775-2	Sequence 2, Appli
14	1029	75.9	245	9	US-09-875-338-23	Sequence 23, Appl
15	952	70.2	290	9	US-10-068-215-23	Sequence 23, Appl
16	952	70.2	290	9	US-09-896-913A-11	Sequence 11, Appl
17	952	70.2	290	10	US-09-794-210-16	Sequence 16, Appl
18	952	70.2	290	10	US-09-910-174A-32	Sequence 32, Appl
19	952	70.2	290	10	US-09-895-837-11	Sequence 11, App
20	952	70.2	290	12	US-10-002-775-11	Sequence 11, Appl
21	796.5	58.7	279	10	US-09-796-858-44	Sequence 44, Appli
22	352	26.0	254	10	US-09-955-866-3	Sequence 3, Appli
23	352	26.0	273	9	US-09-896-913A-2	Sequence 2, Appli
24	352	26.0	273	10	US-09-794-210-2	Sequence 2, Appli
25	352	26.0	273	10	US-09-875-338-15	Sequence 15, Appli
26	352	26.0	273	10	US-09-910-174A-2	Sequence 2, Appli
27	352	26.0	273	10	US-09-955-866-2	Sequence 2, Appli
28	329	24.3	451	10	US-09-895-837-2	Sequence 2, Appli
29	301	22.2	247	9	US-09-875-338-17	Sequence 17, Appl
30	301	22.2	247	9	US-09-896-913A-5	Sequence 5, Appli
31	301	22.2	247	10	US-09-796-858-48	Sequence 48, Appl
32	301	22.2	247	10	US-09-794-210-4	Sequence 4, Appli
33	301	22.2	247	10	US-09-910-174A-31	Sequence 31, Appli
34	301	22.2	247	10	US-09-895-837-5	Sequence 5, Appli
35	292	21.5	534	10	US-09-875-338-11	Sequence 11, Appli
36	292	21.5	534	10	US-09-875-338-7	Sequence 7, Appli
37	289.5	21.3	316	10	US-09-978-295A-13	Sequence 13, Appl
38	289	21.3	316	9	US-09-978-295A-137	Sequence 137, App
39	289	21.3	316	9	US-09-896-697-137	Sequence 137, App
40	289	21.3	316	9	US-09-896-738-14	Sequence 14, Appl
41	289	21.3	316	9	US-09-978-192A-137	Sequence 137, App
42	289	21.3	316	9	US-09-999-832A-137	Sequence 137, App
43	289	21.3	316	10	US-09-790-622-4	Sequence 4, Appli
44	289	21.3	316	9	US-09-978-189-137	Sequence 137, App
45	289	21.3	316	10	US-09-789-561-135	Sequence 135, App
```

ALIGNMENTS

RESULT 1
US-10-068-215-4

Sequence 4, Application US/10068215
Patent No. US20020160000A1
GENERAL INFORMATION:
APPLICANT: Clive Wood
APPLICANT:
APPLICANT: Gordon Freeman
TITLE OF INVENTION: PD-1, A Receptor For B7-4, and Uses Therefor
FILE REFERENCE: GNN-004B
CURRENT APPLICATION NUMBER: US/10/068,215
CURRENT FILING DATE: 2002-02-06
PRIOR APPLICATION NUMBER: 09/645,069
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 60/150,390
PRIOR FILING DATE: 1999-8-23
PRIOR APPLICATION NUMBER: 60/164,897
PRIOR FILING DATE: 1999-11-10
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
LENGTH: 290
TYPE: PRT
ORGANISM: Homo sapiens
US-10-068-215-4

Query Match             100.0%;  Score 1356;  DB 9;  Length 290;
Best Local Similarity   100.0%;  Pred. No. 1e-94;
Matches  261;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  60
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  89

Qy         61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  120
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  149

Qy        121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC  180
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC  209

Qy        181  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV  240

Db   210  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMDV 269
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   241  KKCGIQDTNSKKQSDTHLEET 261
          ||||||||||||||||||||
Db   270  KKCGIQDTNSKKQSDTHLEET 290

RESULT 2
US-09-896-738-12
Sequence 12, Application US/0896738
Patent No. US20020165347A1
GENERAL INFORMATION:
; APPLICANT: Sullivan, John K.
; APPLICANT: Fox, Michael
; TITLE OF INVENTION: B7-Like Molecules and Uses Thereof
; FILE REFERENCE: 00-513-A
; CURRENT APPLICATION NUMBER: US/09,896,738
; "RRENT FILING DATE: 2001-06-29
; IOR APPLICATION NUMBER: 60/215,645
; .RIOR FILING DATE: 2000-06-30
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 12
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-896-738-12

Query Match       100.0%;  Score 1356;  DB 9;  Length 290;
Best Local Similarity   100.0%;  Pred. No. 1e-94;
Matches  261;  Conservative   0;  Mismatches   0;  Indels   0;  Gaps   0;

Qy     1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy    61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMDV 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   210  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMDV 269

Qy   241  KKCGIQDTNSKKQSDTHLEET 261
          ||||||||||||||||||||
Db   270  KKCGIQDTNSKKQSDTHLEET 290

RESULT 3
US-09-896-913A-12
Sequence 12, Application US/0896913A
Patent No. US20020164600A1
GENERAL INFORMATION:
; APPLICANT: Freeman, Gordon
; APPLICANT: Chernova, Irene
; APPLICANT: Chernova, Tatyana
; APPLICANT: Malenkovich, Nelly
; APPLICANT: Wood, Clive
; TITLE OF INVENTION: PD-L2 MOLECULES: NOVEL PD-1 LIGANDS AND
; TITLE OF INVENTION: USES THEREFOR
; FILE REFERENCE: GNN-026A
; CURRENT APPLICATION NUMBER: US/09,896,913A
; CURRENT FILING DATE: 2002-04-15
; PRIOR APPLICATION NUMBER: 60/214,563
; PRIOR FILING DATE: 2000-06-28
; PRIOR APPLICATION NUMBER: 60/270,822

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; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/271,114
; PRIOR FILING DATE: 2001-02-23
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 12
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-896-913A-12

Query Match       100.0%;  Score 1356;  DB 9;  Length 290;
Best Local Similarity   100.0%;  Pred. No. 1e-94;
Matches  261;  Conservative   0;  Mismatches   0;  Indels   0;  Gaps   0;

Qy     1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy    61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMDV 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   210  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMDV 269

Qy   241  KKCGIQDTNSKKQSDTHLEET 261
          ||||||||||||||||||||
Db   270  KKCGIQDTNSKKQSDTHLEET 290

RESULT 4
US-09-915-789A-17
Sequence 17, Application US/0915789A
Patent No. US20020168762A1
GENERAL INFORMATION:
; APPLICANT: Chen, Lieping
; TITLE OF INVENTION: B7-H3 AND B7-H4, NOVEL IMMUNOREGULATORY
; TITLE OF INVENTION: MOLECULES
; FILE REFERENCE: 07039-219001
; CURRENT APPLICATION NUMBER: US/09,915,789A
; CURRENT FILING DATE: 2002-06-04
; PRIOR APPLICATION NUMBER: US 60/220,991
; PRIOR FILING DATE: 2000-07-27
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-915-789A-17

Query Match       100.0%;  Score 1356;  DB 9;  Length 290;
Best Local Similarity   100.0%;  Pred. No. 1e-94;
Matches  261;  Conservative   0;  Mismatches   0;  Indels   0;  Gaps   0;

Qy     1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy    61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

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Qy    181  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV  240
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    210  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV  269

Qy    241  KKCGIQDTNSKKQSDTHLEET  261
           |||||||||||||||||||||
Db    270  KKCGIQDTNSKKQSDTHLEET  290
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RESULT 5
US-09-796-858-42
; Sequence 42, Application US/09796858
; Patent No. US20020051139A1
; GENERAL INFORMATION:
; APPLICANT: Holtzmann, Douglas
; TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING PROGNOSTIC, DIAGNOSTIC,
; TITLE OF INVENTION: PREVENTIVE, THERAPEUTIC, AND OTHER USES
; FILE REFERENCE: 7853-226-999
; CURRENT APPLICATION NUMBER: US/09/796,858
; CURRENT FILING DATE: 2001-03-01
; PRIOR APPLICATION NUMBER: 09/223,094
; PRIOR FILING DATE: 1998-12-30
; PRIOR APPLICATION NUMBER: 09/223,546
; PRIOR FILING DATE: 1998-12-30
; PRIOR APPLICATION NUMBER: 09/224,246
; PRIOR FILING DATE: 1998-12-30
; PRIOR APPLICATION NUMBER: 09/312,359
; PRIOR FILING DATE: 1999-05-14
; PRIOR APPLICATION NUMBER: 09/336,536
; PRIOR FILING DATE: 1999-06-18
; PRIOR APPLICATION NUMBER: 09/342,687
; PRIOR FILING DATE: 1999-06-29
; PRIOR APPLICATION NUMBER: 09/399,723
; PRIOR FILING DATE: 1999-09-20
; PRIOR APPLICATION NUMBER: 09/471,179
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: 09/474,071
; PRIOR FILING DATE: 1999-12-29
; PRIOR APPLICATION NUMBER: 09/474,072
; PRIOR FILING DATE: 1999-12-29
; PRIOR APPLICATION NUMBER: 09/572,002
; PRIOR FILING DATE: 2000-05-14
; PRIOR APPLICATION NUMBER: 09/597,993
; PRIOR FILING DATE: 2000-06-12
; PRIOR APPLICATION NUMBER: 09/599,596
; PRIOR FILING DATE: 2000-06-22
; PRIOR APPLICATION NUMBER: 09/606,565
; PRIOR FILING DATE: 2000-06-29
; PRIOR APPLICATION NUMBER: 09/365,164
; PRIOR FILING DATE: 1999-07-30
; PRIOR APPLICATION NUMBER: 09/630,334
; PRIOR FILING DATE: 2000-07-31
; PRIOR APPLICATION NUMBER: 09/665,666
; PRIOR FILING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 50
; SEQ ID NO 42
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-796-858-42

Query Match   100.0%; Score 1356; DB 10; Length 290;
Best Local Similarity 100.0%; Pred. No. 1e-94;
Matches 261; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149
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RESULT 6
US-09-875-338-2
; Sequence 2, Application US/09875338
; Patent No. US20020050024A1
; GENERAL INFORMATION:
; APPLICANT: MIKESELL, GLEN E.
; APPLICANT: CHANG, HAN
; APPLICANT: FINGER, JOSHUA N.
; APPLICANT: YANG, GUCHEN
; APPLICANT: LU, PIN
; APPLICANT: ZHOU, XIA-DI
; APPLICANT: PEACH, ROBERT
; TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
; TITLE OF INVENTION: IMMUNOMODULATION
; FILE REFERENCE: 3053-4071US2
; CURRENT APPLICATION NUMBER: US/09/875,338
; CURRENT FILING DATE: 2001-06-06
; PRIOR APPLICATION NUMBER: 60/272,107
; PRIOR FILING DATE: 2001-02-28
; PRIOR APPLICATION NUMBER: 60/209,811
; PRIOR FILING DATE: 2000-06-06
; NUMBER OF SEQ ID NOS: 94
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 2
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-875-338-2

Query Match   100.0%; Score 1356; DB 10; Length 290;
Best Local Similarity 100.0%; Pred. No. 1e-94;
Matches 261; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy    1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  89

Qy   61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  149

Qy  121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTNEIFYC  180
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTNEIFYC  209

Qy  181  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV  240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  210  TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV  269

Qy  241  KKCGIQDTNSKKQSDTHLEET  261
          |||||||||||||||||||||
Db  270  KKCGIQDTNSKKQSDTHLEET  290
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RESULT 7
US-09-910-174A-8
; Sequence 8, Application US/09910174A
; Patent No. US20020106730A1
; GENERAL INFORMATION:
; APPLICANT: Coyle, Anthony J.

; APPLICANT: Fraser, Christopher C.
; APPLICANT: Manning, Stephen
; TITLE OF INVENTION: B7-H2 Molecules, No. US20020106730A1e1 Members of the B7
; TITLE OF INVENTION: Family and Uses Thereof
; FILE REFERENCE: 35800/236924
; CURRENT APPLICATION NUMBER: US/09/910,174A
; CURRENT FILING DATE: 2001-07-20
; PRIOR APPLICATION NUMBER: US 09/620,461
; PRIOR FILING DATE: 2000-07-20
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-910-174A-8

Query Match   100.0%;   Score 1356;   DB 10;   Length 290;
Best Local Similarity 100.0%;   Pred. No. 1e-94;
Matches 261;   Conservative 0;   Mismatches 0;   Indels 0;   Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 240
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 210 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 269

Qy 241 KKCGIQDTNSKKQSDTHLEET 261
         |||||||||||||||||||||
Db 270 KKCGIQDTNSKKQSDTHLEET 290

RESULT 8
US-09-955-866-6
Sequence 6, Application US/09955866
Patent No. US20020107363A1
GENERAL INFORMATION:
; APPLICANT: Fox, Michael
; APPLICANT: Sullivan, John K.
; APPLICANT: Holst, Paige
; APPLICANT: Yoshinaga, Steven Kiyoshi
; TITLE OF INVENTION: B7-Like Polypeptides and Uses Thereof
; FILE REFERENCE: 00,759-A
; CURRENT APPLICATION NUMBER: US/09/955,866
; CURRENT FILING DATE: 2001-09-19
; PRIOR APPLICATION NUMBER: 60/233,867
; PRIOR FILING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 6
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-955-866-6

Query Match   100.0%;   Score 1356;   DB 10;   Length 290;
Best Local Similarity 100.0%;   Pred. No. 1e-94;
Matches 261;   Conservative 0;   Mismatches 0;   Indels 0;   Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 240
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 210 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 269

Qy 241 KKCGIQDTNSKKQSDTHLEET 261
         |||||||||||||||||||||
Db 270 KKCGIQDTNSKKQSDTHLEET 290

RESULT 9
US-09-895-837-12
Sequence 12, Application US/09895837
Patent No. US20020110836A1
GENERAL INFORMATION:
; APPLICANT: Freeman, Gordon
; APPLICANT: Chernova, Irene
; APPLICANT: Chernova, Tatyana
; APPLICANT: Malenkovich, Nelly
; APPLICANT: Wood, Clive
; APPLICANT: Latchman, Yvette
; APPLICANT: Sharpe, Arlene H.
; TITLE OF INVENTION: PD-L2 MOLECULES: NOVEL PD-1 LIGANDS AND
; TITLE OF INVENTION: USES THEREFOR
; FILE REFERENCE: GNN-026B
; CURRENT APPLICATION NUMBER: US/09/895,837
; CURRENT FILING DATE: 2001-06-28
; PRIOR APPLICATION NUMBER: 60/214,563
; PRIOR FILING DATE: 2000-06-28
; PRIOR APPLICATION NUMBER: 60/270,822
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/271,114
; PRIOR FILING DATE: 2001-02-23
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 12
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-895-837-12

Query Match   100.0%;   Score 1356;   DB 10;   Length 290;
Best Local Similarity 100.0%;   Pred. No. 1e-94;
Matches 261;   Conservative 0;   Mismatches 0;   Indels 0;   Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 240
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 210 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 269

Qy 241 KKCGIQDTNSKKQSDTHLEET 261
         |||||||||||||||||||||
Db 270 KKCGIQDTNSKKQSDTHLEET 290

RESULT 10
US-10-002-775-4
; Sequence 4, Application US/10002775
; Patent No. US20020102651A1
; GENERAL INFORMATION:
; APPLICANT: Gordon Freeman
; APPLICANT: Vassiliki Boussiotis
; APPLICANT: Tatyana Chernova
; APPLICANT: Nelly Malenkovich
; TITLE OF INVENTION: NOVEL B7-4 MOLECULES AND USES THEREFOR
; FILE REFERENCE: GNN-004ADV
; CURRENT APPLICATION NUMBER: US/10/002,775
; CURRENT FILING DATE: 2001-11-02
; PRIOR APPLICATION NUMBER: US 09/644,934
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-08-23
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: Patentin Ver. 2.0
;
; SEQ ID NO 4
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens

US-10-002-775-4

Query Match   100.0%; Score 1356; DB 12; Length 290;
Best Local Similarity 100.0%; Pred. No. 1e-94;
Matches 261; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 210 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKGRMMDV 269

Qy 241 KKCGIQDTNSKKQSDTHLEET 261
        |||||||||||||||||||||
Db 270 KKCGIQDTNSKKQSDTHLEET 290

RESULT 11
US-09-915-789A-23
; Sequence 23, Application US/09915789A
; Patent No. US20020168762A1
; GENERAL INFORMATION:
; APPLICANT: Chen, Lieping
; TITLE OF INVENTION: B7-H3 AND B7-H4, NOVEL IMMUNOREGULATORY
; TITLE OF INVENTION: MOLECULES
; FILE REFERENCE: 07039-219001
; CURRENT APPLICATION NUMBER: US/09/915,789A
; CURRENT FILING DATE: 2002-06-04
; PRIOR APPLICATION NUMBER: US 60/220,991
; PRIOR FILING DATE: 2000-07-27
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSEQ for Windows Version 4.0
;
; SEQ ID NO 23
; LENGTH: 220
; TYPE: PRT
; ORGANISM: Homo sapiens

US-09-915-789A-23

Query Match   81.4%; Score 1104; DB 9; Length 220;
Best Local Similarity 100.0%; Pred. No. 5.4e-76;

Matches 211; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   10 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 69

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   70 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 129

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 130 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 189

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERTH 211
        ||||||||||||||||||||||||||||||
Db 190 TFRRLDPEENHTAELVIPELPLAHPPNERTH 220

RESULT 12
US-09-875-338-5
; Sequence 5, Application US/09875338
; Patent No. US20020095024A1
; GENERAL INFORMATION:
; APPLICANT: MIKESELL, GLEN E.
; APPLICANT: CHANG, HAN
; APPLICANT: FINGER, JOSHUA N.
; APPLICANT: YANG, GUCHEN
; APPLICANT: LU, PIN
; APPLICANT: ZHOU, XIA-DI
; APPLICANT: PEACH, ROBERT
; TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
; TITLE OF INVENTION: IMMUNOMODULATION
; FILE REFERENCE: 3053-4071US2
; CURRENT APPLICATION NUMBER: US/09/875,338
; CURRENT FILING DATE: 2001-06-06
; PRIOR APPLICATION NUMBER: 60/272,107
; PRIOR FILING DATE: 2001-02-28
; PRIOR APPLICATION NUMBER: 60/209,811
; PRIOR FILING DATE: 2000-06-06
; NUMBER OF SEQ ID NOS: 94
; SOFTWARE: Patentin Ver. 2.1
;
; SEQ ID NO 5
; LENGTH: 480
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: fusion construct

US-09-875-338-5

Query Match   80.8%; Score 1096; DB 10; Length 480;
Best Local Similarity 100.0%; Pred. No. 5.4e-75;
Matches 210; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   34 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 93

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   94 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 153

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 154 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 213

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
        |||||||||||||||||||||||||||||
Db 214 TFRRLDPEENHTAELVIPELPLAHPPNERT 243

RESULT 13

US-10-068-215-2
; Sequence 2, Application US/10068215
; Patent No. US20020160000A1
; GENERAL INFORMATION:
; APPLICANT: Clive Wood
; TITLE OF INVENTION: PD-1, A Receptor For B7-4, and Uses Therefor
; FILE REFERENCE: GNN-004B
; CURRENT APPLICATION NUMBER: US/10/068,215
; CURRENT FILING DATE: 2002-02-06
; PRIOR APPLICATION NUMBER: 09,645,069
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-8-23
; PRIOR APPLICATION NUMBER: 60/164,897
; PRIOR FILING DATE: 1999-11-10
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2
; LENGTH: 245
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-068-215-2

Query Match 75.9%; Score 1029; DB 9; Length 245;
Best Local Similarity 100.0%; Pred. No. 2.5e-70;
Matches 198; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy  121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy  181 TFRRLDPEENHTAELVIP 198
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Db  210 TFRRLDPEENHTAELVIP 227
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RESULT 14
US-10-002-775-2
; 'ence 2, Application US/10002775
; .nt No. US20020102651A1
; G__ERAL INFORMATION:
; APPLICANT: Gordon Freeman
; APPLICANT: Vassiliki Boussiotis
; APPLICANT: Tatyana Chernova
; APPLICANT: Nelly Malenkovich
; TITLE OF INVENTION: NOVEL B7-4 MOLECULES AND USES THEREFOR
; FILE REFERENCE: GNN-004ADV
; CURRENT APPLICATION NUMBER: US/10/002,775
; CURRENT FILING DATE: 2001-11-02
; PRIOR APPLICATION NUMBER: US 09/644,934
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-08-23
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2
; LENGTH: 245
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-002-775-2

Query Match 75.9%; Score 1029; DB 12; Length 245;
Best Local Similarity 100.0%; Pred. No. 2.5e-70;
Matches 198; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy    1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  60
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Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy  121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy  181 TFRRLDPEENHTAELVIP 198
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Db  210 TFRRLDPEENHTAELVIP 227
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RESULT 15
US-10-068-215-23
; Sequence 23, Application US/10068215
; Patent No. US20020160000A1
; GENERAL INFORMATION:
; APPLICANT: Clive Wood
; TITLE OF INVENTION: PD-1, A Receptor For B7-4, and Uses Therefor
; FILE REFERENCE: GNN-004B
; CURRENT APPLICATION NUMBER: US/10/068,215
; CURRENT FILING DATE: 2002-02-06
; PRIOR APPLICATION NUMBER: 09/645,069
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-8-23
; PRIOR APPLICATION NUMBER: 60/164,897
; PRIOR FILING DATE: 1999-11-10
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 23
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-068-215-23

Query Match 70.2%; Score 952; DB 9; Length 290;
Best Local Similarity 69.5%; Pred. No. 1.8e-64;
Matches 182; Conservative 33; Mismatches 45; Indels 2; Gaps 2;

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Qy    1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK  60
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Db   30 VEYGSNVTMECRFPVERELDLLALVVYWEKEDEQVIQFVAGEEDLKPQHSNFRGRASLPK  89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
        |||::|||||||||||||||||||:|:||||||||||:|||||||:||||||||-:||:||
Db   90 DQLLKGNAALQITDVKLQDAGVYCCIISYGGADYKRITLKVNAPYRKINQRI-SVPDATS 148

Qy  121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
        |||||||||||:|||||||:|||||:|||:::|:|:::|::|||||:|||:||::|:||||
Db  149 EHELTCQAEGYPEAEVIWTNSDHQPVSGKRSVTTSRTEGMLLNVTSSLRVNATANDVFYC 208

Qy  181 TFRRLDPEENHTAELVIPELPLAHPPNERTHLVILGAILLCLGVALTFIFRLRKG-RMMD 239
        ||:|::|:||:|||:||||||:|||||:||:|:||:|||::|:|:::|:||||| ||::D
Db  209 TFWRSQPGQNHTAELIIPELPATHPPQNRHWVLLGSILLFLIVVSTVLLFLRKQVRMLD 268

Qy  240 VKKCCIQDDTNSKKQSDTHLEET 261
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Db  269 VEKCCGVEDTSSKNRNDTQFEET 290
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Search completed: January 12, 2003, 10:19:07
Job time : 11.5287 secs

OM protein - protein search, using sw model

GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.

Run on: January 12, 2003, 10:10:43 ; Search time 8.47134 Seconds
                                                  (without alignments)
                                                  480.942 Million cell updates/sec

Title: US-09-649-108-10
Perfect score: 1096
Sequence: 1 VEYGSNMTIECKFPVEKQLD...........HTAELVIPELPLAHPPNERT 210

Scoring table: BLOSUM62
                     Gapop 10.0 , Gapext 0.5

Searched: 118974 seqs, 19401057 residues

All number of hits satisfying chosen parameters: 118974

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-Processing: Minimum Match 0%
                     Maximum Match 100%
                     Listing first 45 summaries

Database :   Published_Applications_AA:*
                     1: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
                     2: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
                     3: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
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                     5: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
                     6: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
                     7: /cgn2_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*
                     8: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
                     9: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
                     10: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
                     11: /cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
                     12: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
                     13: /cgn2_6/ptodata/2/pubpaa/US60_NEW_PUB.pep:*
                     14: /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Match Length	DB	ID	Description
1	1096	100.0	220	9	US-09-915-789A-23	Sequence 23, Appl
2	1096	100.0	290	9	US-10-068-215-4	Sequence 4, Appli
3	1096	100.0	290	9	US-09-896-738-12	Sequence 12, Appl
4	1096	100.0	290	9	US-09-896-913A-12	Sequence 12, Appl
5	1096	100.0	290	9	US-09-915-789A-17	Sequence 17, Appl
6	1096	100.0	290	10	US-09-796-858-42	Sequence 42, Appl
7	1096	100.0	290	10	US-09-875-338-2	Sequence 2, Appli
8	1096	100.0	290	10	US-09-910-174A-8	Sequence 8, Appli
9	1096	100.0	290	10	US-09-955-866-6	Sequence 6, Appli
10	1096	100.0	290	12	US-09-895-837-12	Sequence 12, Appl
10	1096	100.0	480	12	US-09-002-775-4	Sequence 4, Appli
11	1096	100.0	245	9	US-10-068-215-2	Sequence 2, Appli
12	1029	93.9	245	12	US-09-875-338-5	Sequence 5, Appli
13	1029	93.9	245	9	US-10-002-775-2	Sequence 2, Appli
14	1096	100.0	290	9	US-10-068-215-23	Sequence 23, Appl
15	815.5	74.4	290	9	US-09-896-913A-11	Sequence 11, Appl
16	815.5	74.4	290	10	US-09-794-210-16	Sequence 16, Appl
17	815.5	74.4	290	9	US-09-910-174A-31	Sequence 32, Appl
18	815.5	74.4	290	10	US-09-895-837-11	Sequence 11, App
19	815.5	74.4	290	10	US-09-895-837-11	Sequence 11, App
20	815.5	74.4	290	12	US-10-002-775-11	Sequence 11, Appl
21	697	63.6	279	10	US-09-796-858-44	Sequence 44, Appl
22	336	30.7	254	10	US-09-955-866-3	Sequence 3, Appli
23	336	30.7	273	9	US-09-896-913A-2	Sequence 2, Appli
24	336	30.7	273	10	US-09-794-210-2	Sequence 2, Appli
25	336	30.7	273	10	US-09-875-338-15	Sequence 15, Appl
26	336	30.7	273	10	US-09-910-174A-2	Sequence 2, Appli
27	336	30.7	273	10	US-09-955-866-2	Sequence 2, Appli
28	336	30.7	273	12	US-09-895-837-2	Sequence 2, Appli
29	329	30.0	451	10	US-09-875-333-17	Sequence 17, Appl
30	289.5	26.4	247	9	US-09-896-913A-5	Sequence 5, Appl
31	289.5	26.4	247	10	US-09-796-858-48	Sequence 48, Appl
32	289.5	26.4	247	10	US-09-794-210-4	Sequence 4, Appli
33	289.5	26.4	247	9	US-09-910-174A-31	Sequence 31, Appl
34	289.5	26.4	247	10	US-09-895-837-5	Sequence 5, Appli
35	265.5	24.2	316	11	US-09-875-338-11	Sequence 11, Appl
36	265.5	24.2	534	10	US-09-875-338-7	Sequence 7, Appl
37	265.5	24.2	698	10	US-09-875-338-9	Sequence 9, Appli
38	262.5	24.0	316	9	US-09-978-2955-137	Sequence 137, App
39	262.5	24.0	316	9	US-09-896-738-14	Sequence 14, App
40	262.5	24.0	316	9	US-09-978-697-137	Sequence 137, App
41	262.5	24.0	316	10	US-09-999-1924-137	Sequence 137, App
42	262.5	24.0	316	9	US-09-999-832A-137	Sequence 137, App
43	262.5	24.0	316	9	US-09-790-622-4	Sequence 4, Appli
44	262.5	24.0	316	9	US-09-976-189-137	Sequence 137, App
45	262.5	24.0	316	10	US-09-789-561-135	Sequence 135, App

ALIGNMENTS

RESULT 1
US-09-915-789A-23

Sequence 23, Application US/0991578946

Patent No. US20020168762A1

GENERAL INFORMATION:

APPLICANT: Chen, Lieping
TITLE OF INVENTION: B7-H3 AND B7-H4, NOVEL IMMUNOREGULATORY
TITLE OF INVENTION: MOLECULES
FILE REFERENCE: 07039-219001
CURRENT APPLICATION NUMBER: US/09/915,789A
CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US 60/220,991
PRIOR FILING DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 23
LENGTH: 220
TYPE: PRT
ORGANISM: Homo sapiens
US-09-915-789A-23

Query Match   100.0%; Score 1096; DB 9; Length 220;
Best Local Similarity   100.0%;
Matches 210; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
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Db   10 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 69

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKIINQRILVVDPVTS 120
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   70 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKIINQRILVVDPVTS 129

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   130 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 189

Qy   181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
             |||||||||||||||||||||||||||||
Db   190 TFRRLDPEENHTAELVIPELPLAHPPNERT 219

RESULT 2
US-10-068-215-4
; Sequence 4, Application US/10068215
; Patent No. US20020160000A1
; GENERAL INFORMATION:
; APPLICANT: Clive Wood
; APPLICANT: Gordon Freeman
; TITLE OF INVENTION: PD-1, A Receptor For B7-4, and Uses Therefor
; FILE REFERENCE: GNN-004B
; CURRENT APPLICATION NUMBER: US/10/068,215
; CURRENT FILING DATE: 2002-02-06
; PRIOR APPLICATION NUMBER: 09/645,069
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-8-23
; PRIOR APPLICATION NUMBER: 60/164,897
; PRIOR FILING DATE: 1999-11-10
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.0
; .ID NO 4
; .ENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-068-215-4

Query Match   100.0%;   Score 1096; DB 9; Length 290;
Best Local Similarity   100.0%;   Pred. No. 3e-73;
Matches 210; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy    1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   60
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   89

Qy   61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  120
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  149

Qy  121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC  180
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC  209

Qy  181  TFRRLDPEENHTAELVIPELPLAHPPNERT  210
         |||||||||||||||||||||||||||||
Db  210  TFRRLDPEENHTAELVIPELPLAHPPNERT  239
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RESULT 3
US-?-896-738-12
; ...ent No. US20020153347A1
; ...ience 12, Application US/0896738
; GENERAL INFORMATION:
; APPLICANT: Fox, Michael
; APPLICANT: Sullivan, John K.
; APPLICANT: Fang, Mei
; TITLE OF INVENTION: B7-Like Molecules and Uses Thereof
; FILE REFERENCE: 00-513-A
; CURRENT APPLICATION NUMBER: US/09/896,738
; CURRENT FILING DATE: 2001-06-29
; PRIOR APPLICATION NUMBER: 60/215,645
; PRIOR FILING DATE: 2000-06-30
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 12
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-896-738-12

Query Match   100.0%;   Score 1096; DB 9; Length 290;
Best Local Similarity   100.0%;   Pred. No. 3e-73;
Matches 210; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy    1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   60
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RESULT 4
US-09-896-913A-12
; Sequence 12, Application US/0896913A
; Patent No. US20020164600A1
; GENERAL INFORMATION:
; APPLICANT: Freeman, Gordon
; APPLICANT: Chernova, Irene
; APPLICANT: Chernova, Tatyana
; APPLICANT: Malenkovich, Nelly
; APPLICANT: Wood, Clive
; TITLE OF INVENTION: PD-L2 MOLECULES: NOVEL PD-1 LIGANDS AND
; TITLE OF INVENTION: USES THEREFOR
; FILE REFERENCE: GNN-026A
; CURRENT APPLICATION NUMBER: US/09/896,913A
; CURRENT FILING DATE: 2002-04-15
; PRIOR APPLICATION NUMBER: 60/214,563
; PRIOR FILING DATE: 2000-06-28
; PRIOR APPLICATION NUMBER: 60/270,822
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/271,114
; PRIOR FILING DATE: 2001-02-23
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 12
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-896-913A-12

Query Match   100.0%;   Score 1096; DB 9; Length 290;
Best Local Similarity   100.0%;   Pred. No. 3e-73;
Matches 210; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy    1  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   60
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30  VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK   89

Qy   61  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  120
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90  DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS  149

Qy  121  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC  180
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  150  EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC  209

Qy  181  TFRRLDPEENHTAELVIPELPLAHPPNERT  210
         |||||||||||||||||||||||||||||
Db  210  TFRRLDPEENHTAELVIPELPLAHPPNERT  239
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RESULT 5
US-09-915-789A-17
; Sequence 17, Application US/0915789A
; Patent No. US20020168762A1
; GENERAL INFORMATION:
; APPLICANT: Chen, Lieping
; TITLE OF INVENTION: B7-H3 AND B7-H4, NOVEL IMMUNOREGULATORY

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; TITLE OF INVENTION: MOLECULES
; FILE REFERENCE: 07039-219001
; CURRENT APPLICATION NUMBER: US/09/915,789A
; CURRENT FILING DATE: 2002-06-04
; PRIOR APPLICATION NUMBER: US 60/220,991
; PRIOR FILING DATE: 2000-07-27
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-915-789A-17

  Query Match           100.0%;  Score 1096;  DB 9;   Length 290;
  Best Local Similarity 100.0%;  Pred. No. 3e-73;
  Matches 210;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy  61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRLLVVDPVTS 120
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRLLVVDPVTS 149

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
       ||||||||||||||||||||||||||||||
Db 210 TFRRLDPEENHTAELVIPELPLAHPPNERT 239
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```
RESULT 6
US-09-796-858-42
Sequence 42, Application US/09796858
Patent No. US20020051139A1
GENERAL INFORMATION:
; APPLICANT: Holtzmann, Douglas
; TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING PROGNOSTIC, DIAGNOSTIC,
; TITLE OF INVENTION: PREVENTIVE, THERAPEUTIC, AND OTHER USES
; FILE REFERENCE: 7853-226-999
; CURRENT APPLICATION NUMBER: US/09/796,858
; CURRENT FILING DATE: 2001-03-01
; PRIOR APPLICATION NUMBER: 09/223,094
; PRIOR FILING DATE: 1998-12-30
; PRIOR APPLICATION NUMBER: 09/223,546
; PRIOR FILING DATE: 1998-12-30
; PRIOR APPLICATION NUMBER: 09/224,246
; PRIOR FILING DATE: 1998-12-30
; PRIOR APPLICATION NUMBER: 09/312,359
; PRIOR FILING DATE: 1999-05-14
; PRIOR APPLICATION NUMBER: 09/336,536
; PRIOR FILING DATE: 1999-06-18
; PRIOR APPLICATION NUMBER: 09/342,687
; PRIOR FILING DATE: 1999-06-29
; PRIOR APPLICATION NUMBER: 09/399,723
; PRIOR FILING DATE: 1999-09-20
; PRIOR APPLICATION NUMBER: 09/471,179
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: 09/474,071
; PRIOR FILING DATE: 1999-12-29
; PRIOR APPLICATION NUMBER: 09/474,072
; PRIOR FILING DATE: 1999-12-29
; PRIOR APPLICATION NUMBER: 09/597,993
; PRIOR FILING DATE: 2000-06-12
; PRIOR APPLICATION NUMBER: 09/599,596
; PRIOR FILING DATE: 2000-06-22
; PRIOR APPLICATION NUMBER: 09/606,565
; PRIOR FILING DATE: 2000-06-29
; PRIOR APPLICATION NUMBER: 09/365,164
; PRIOR FILING DATE: 1999-07-30
; PRIOR APPLICATION NUMBER: 09/630,334
; PRIOR FILING DATE: 2000-07-31
; PRIOR APPLICATION NUMBER: 09/665,666
; PRIOR FILING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 50
; SEQ ID NO 42
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-796-858-42

  Query Match           100.0%;  Score 1096;  DB 9;   Length 290;
  Best Local Similarity 100.0%;  Pred. No. 3e-73;
  Matches 210;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy  61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRLLVVDPVTS 120
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRLLVVDPVTS 149

Qy 121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy 181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
       ||||||||||||||||||||||||||||||
Db 210 TFRRLDPEENHTAELVIPELPLAHPPNERT 239
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RESULT 7
US-09-875-338-2
Sequence 2, Application US/09875338
Patent No. US20020095024A1
GENERAL INFORMATION:
; APPLICANT: MIKESELL, GLEN E.
; APPLICANT: CHANG, HAN
; APPLICANT: FINGER, JOSHUA N.
; APPLICANT: YANG, GUCHEN
; APPLICANT: LU, PIN
; APPLICANT: ZHOU, XIA-DI
; APPLICANT: PEACH, ROBERT
; TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
; TITLE OF INVENTION: IMMUNOMODULATION
; FILE REFERENCE: 3053-4071US2
; CURRENT APPLICATION NUMBER: US/09/875,338
; CURRENT FILING DATE: 2001-06-06
; PRIOR APPLICATION NUMBER: 60/272,107
; PRIOR FILING DATE: 2001-02-28
; PRIOR APPLICATION NUMBER: 60/209,811
; PRIOR FILING DATE: 2000-06-06
; NUMBER OF SEQ ID NOS: 94
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-875-338-2

  Query Match           100.0%;  Score 1096;  DB 10;   Length 290;
  Best Local Similarity 100.0%;  Pred. No. 3e-73;
  Matches 210;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
       ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy  61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRLLVVDPVTS 120
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RESULT 8
US-09-910-174A-8

Sequence 8, Application US/0991011074A
Patent No. US20020106730A1
GENERAL INFORMATION:
 APPLICANT: Coyle, Anthony J.
 APPLICANT: Fraser, Christopher C.
 APPLICANT: Manning, Stephen
 TITLE OF INVENTION: B7-H2 Molecules, Members of the B7
 Family and Uses Thereof
 FILE REFERENCE: 35800/236924
 CURRENT APPLICATION NUMBER: US/09/910,174A
 CURRENT FILING DATE: 2001-07-20
 PRIOR APPLICATION NUMBER: US 09/620,461
 PRIOR FILING DATE: 2000-07-20
 NUMBER OF SEQ ID NOS: 32
 SOFTWARE: FastSEQ for Windows Version 4.0
 SEQ ID NO 8
 LENGTH: 290
 TYPE: PRT
 ORGANISM: Homo sapiens
US-09-910-174A-8

Query Match   100.0%; Score 1096; DB 10; Length 290;
Best Local Similarity   100.0%; Pred. No. 3e-73;
Matches 210; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
           |||||||||||||||||||||||||||||||
Db   210 TFRRLDPEENHTAELVIPELPLAHPPNERT 239

RESULT 9
US-09-955-866-6

Sequence 6, Application US/09955866
Patent No. US20020107363A1
GENERAL INFORMATION:
 APPLICANT: Fox, Michael
 APPLICANT: Sullivan, John K.
 APPLICANT: Holst, Paige
 APPLICANT: Yoshinaga, Steven Kiyoshi
 TITLE OF INVENTION: B7-Like Polypeptides and Uses Thereof
 FILE REFERENCE: 00.759-A
 CURRENT APPLICATION NUMBER: US/09/955,866
 CURRENT FILING DATE: 2001-09-19
 PRIOR APPLICATION NUMBER: 60/233,867
 PRIOR FILING DATE: 2000-09-20
 NUMBER OF SEQ ID NOS: 30
 SOFTWARE: PatentIn Ver. 2.0

 SEQ ID NO 6
 LENGTH: 290
 TYPE: PRT
 ORGANISM: Homo sapiens
US-09-955-866-6

Query Match   100.0%; Score 1096; DB 10; Length 290;
Best Local Similarity   100.0%; Pred. No. 3e-73;
Matches 210; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
           |||||||||||||||||||||||||||||||
Db   210 TFRRLDPEENHTAELVIPELPLAHPPNERT 239

RESULT 10
US-09-895-837-12

Sequence 12, Application US/09895837
Patent No. US20020110836A1
GENERAL INFORMATION:
 APPLICANT: Freeman, Gordon
 APPLICANT: Chernova, Irene
 APPLICANT: Chernova, Tatyana
 APPLICANT: Malenkovich, Nelly
 APPLICANT: Wood, Clive
 APPLICANT: Latchman, Yvette
 APPLICANT: Sharpe, Arlene H.
 TITLE OF INVENTION: PD-L2 MOLECULES: NOVEL PD-1 LIGANDS AND
 USES THEREFOR
 FILE REFERENCE: GNN-026B
 CURRENT APPLICATION NUMBER: US/09/895,837
 CURRENT FILING DATE: 2001-06-28
 PRIOR APPLICATION NUMBER: 60/214,563
 PRIOR FILING DATE: 2000-06-28
 PRIOR APPLICATION NUMBER: 60/270,822
 PRIOR FILING DATE: 2001-02-23
 PRIOR APPLICATION NUMBER: 60/271,114
 PRIOR FILING DATE: 2001-02-23
 NUMBER OF SEQ ID NOS: 12
 SOFTWARE: FastSEQ for Windows Version 4.0
 SEQ ID NO 12
 LENGTH: 290
 TYPE: PRT
 ORGANISM: Homo sapiens
US-09-895-837-12

Query Match   100.0%; Score 1096; DB 10; Length 290;
Best Local Similarity   100.0%; Pred. No. 3e-73;
Matches 210; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
           |||||||||||||||||||||||||||||||
Db   210 TFRRLDPEENHTAELVIPELPLAHPPNERT 239

RESULT 11
US-10-002-775-4
; Sequence 4, Application US/10002775
; Patent No. US20020102651A1
; CURRENT APPLICATION NUMBER: US/10,002,775
; CURRENT FILING DATE: 2001-11-02
; PRIOR APPLICATION NUMBER: US 09/644,934
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-08-23
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 4
; LENGTH: 290
; TYPE: PRT
; ORGANISM: Homo sapiens
; GENERAL INFORMATION:
; APPLICANT: Nelly Malenkovich
; APPLICANT: Tatyana Chernova
; APPLICANT: Vassiliki Boussiotis
; APPLICANT: Gordon Freeman
; TITLE OF INVENTION: NOVEL B7-4 MOLECULES AND USES THEREFOR
; FILE REFERENCE: GNN-004ADV
; CURRENT APPLICATION NUMBER: US/10,002,775

Query Match   100.0%; Score 1096; DB 12; Length 290;
Best Local Similarity   100.0%;
Matches 210; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
          |||||||||||||||||||||||||||||
Db   210 TFRRLDPEENHTAELVIPELPLAHPPNERT 239

RESULT 12
US-09-875-338-5
; Sequence 5, Application US/0987538
; Patent No. US20020095024A1
; GENERAL INFORMATION:
; APPLICANT: MIKESELL, GLEN E.
; APPLICANT: CHANG, HAN
; APPLICANT: FINGER, JOSHUA N.
; APPLICANT: YANG, GUCHEN
; APPLICANT: LU, PIN
; APPLICANT: ZHOU, XIA-DI
; APPLICANT: PEACH, ROBERT
; TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND POLYPEPTIDES USEFUL FOR
; TITLE OF INVENTION: IMMUNOMODULATION
; FILE REFERENCE: 3053-4071US2
; CURRENT APPLICATION NUMBER: US/09/875,338
; CURRENT FILING DATE: 2001-06-06
; PRIOR APPLICATION NUMBER: 60/272,107
; PRIOR FILING DATE: 2001-02-28
; PRIOR APPLICATION NUMBER: 60/209,811
; PRIOR FILING DATE: 2000-06-06

; NUMBER OF SEQ ID NOS: 94
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 5
; LENGTH: 480
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: fusion construct
US-09-875-338-5

Query Match   100.0%; Score 1096; DB 10; Length 480;
Best Local Similarity   100.0%;
Matches 210; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   34 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 93

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   94 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 153

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   154 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 213

Qy   181 TFRRLDPEENHTAELVIPELPLAHPPNERT 210
          |||||||||||||||||||||||||||||
Db   214 TFRRLDPEENHTAELVIPELPLAHPPNERT 243

RESULT 13
US-10-068-215-2
; Sequence 2, Application US/10068215
; Patent No. US20020160000A1
; GENERAL INFORMATION:
; APPLICANT: Clive Wood
; APPLICANT: Gordon Freeman
; TITLE OF INVENTION: PD-1, A Receptor For B7-4, and Uses Therefor
; FILE REFERENCE: GNN-004B
; CURRENT APPLICATION NUMBER: US/10/068,215
; CURRENT FILING DATE: 2002-02-06
; PRIOR APPLICATION NUMBER: 09/645,069
; PRIOR FILING DATE: 2000-08-23
; PRIOR APPLICATION NUMBER: 60/150,390
; PRIOR FILING DATE: 1999-8-23
; PRIOR APPLICATION NUMBER: 60/164,897
; PRIOR FILING DATE: 1999-11-10
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2
; LENGTH: 245
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-068-215-2

Query Match   93.9%; Score 1029; DB 9; Length 245;
Best Local Similarity   100.0%;
Matches 198; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy   61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

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Qy   181 TFFRLDPEENHTAELVIP 198
         ||||||||||||||||||
Db   210 TFFRLDPEENHTAELVIP 227
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RESULT 14
US-10-002-775-2
Sequence 2, Application US/10002775
Patent No. US20020102651A1
GENERAL INFORMATION:
APPLICANT: Gordon Freeman
APPLICANT: Vassiliki Boussiotis
APPLICANT: Tatyana Chernova
APPLICANT: Nelly Malenkovich
TITLE OF INVENTION: NOVEL B7-4 MOLECULES AND USES THEREFOR
FILE REFERENCE: GNN-004ADV
CURRENT APPLICATION NUMBER: US/10/002,775
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: US 09/644,934
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 60/150,390
PRIOR FILING DATE: 1999-08-23
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 245
TYPE: PRT
ORGANISM: Homo sapiens
US-10-002-775-2

Query Match   93.9%; Score 1029; DB 12; Length 245;
Best Local Similarity   100.0%; Pred. No. 1.9e-68;
Matches 198; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

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Qy     1 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 60
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Db    30 VEYGSNMTIECKFPVEKQLDLAALIVYWEMEDKNIIQFVHGEEDLKVQHSSYRQRARLLK 89

Qy    61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    90 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 149

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   150 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 209

Qy   181 TFFRLDPEENHTAELVIP 198
         ||||||||||||||||||
Db   210 TFFRLDPEENHTAELVIP 227
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RESULT 15
US-10-068-215-23
Sequence 23, Application US/10068215
Patent No. US20020160000A1
GENERAL INFORMATION:
APPLICANT: Clive Wood
APPLICANT: Gordon Freeman
TITLE OF INVENTION: PD-1, A Receptor For B7-4, and Uses Therefor
FILE REFERENCE: GNN-004B
CURRENT APPLICATION NUMBER: US/10/068,215
CURRENT FILING DATE: 2002-02-06
PRIOR APPLICATION NUMBER: 09/645,069
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 60/150,390
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 60/150,390
PRIOR FILING DATE: 1999-8-23
PRIOR APPLICATION NUMBER: 60/164,897
PRIOR FILING DATE: 1999-11-10
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 23
LENGTH: 290
TYPE: PRT

ORGANISM: Mus musculus
US-10-068-215-23

Query Match   74.4%; Score 815.5; DB 9; Length 290;
Best Local Similarity   72.9%; Pred. No. 8.4e-53;
Matches 153; Conservative 23; Mismatches 33; Indels   1; Gaps   1;

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         ||||||:||:|:||||:||||||:|||| |||||||||||||||| |||:||:||||| ||
Db    30 VEYGSNVTMECRFPVERELDLLALVVYWEKEDEQVIQFVAGEEDLKPQHSNFRGRASLPK 89

Qy    61 DQLSLGNAALQITDVKLQDAGVYRCMISYGGADYKRITVKVNAPYNKINQRILVVDPVTS 120
         || |||||||||||||||||||| :|||||||:||||| |||||| |:|||| |||
Db    90 DQLLKGNAALQITDVKLQDAGVYCCIISYGGADYKRITLKVNAPYRKINQRI-SVDPATS 148

Qy   121 EHELTCQAEGYPKAEVIWTSSDHQVLSGKTTTTNSKREEKLFNVTSTLRINTTTNEIFYC 180
         |||:||||||:||||||||.|:||:| |:| | ||| :.. :..||||| |||:|:.|:|||||
Db   149 EHELICQAEGYPEAEVIWTNSDHQPVSGKRSVTTSRIEGMLLNVTSSLRVNATANDVFYC 208

Qy   181 TFFRLDPEENHTAELVIPELPLAHPPNERT 210
         |||||||||:|||||||||:|||:| ||
Db   209 TFWRSQPGQNHTAELIIPELPATHPPQNRT 238
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Search completed: January 12, 2003, 10:19:08
Job time : 9.47134 secs